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IN THE UNITED STATES PATENT AND TRADEMARK OFFICE

APPLICANTS:

Guo, et al.

ASSIGNEE:

CURAGEN CORP.

SERIAL NUMBER:

10/074,978

EXAMINER:

Not Yet Assigned

FILING DATE:

February 12, 2002

ART UNIT:

1646

For:

NOVEL HUMAN PROTEINS AND NUCLEIC ACIDS ENCODING

SAME

Box Sequence Commissioner For Patents Washington, D.C. 20231

STATEMENT IN SUPPORT OF COMPUTER READA FORM SUBMISSION UNDER 37 C.F.R. § 1.821(f)

I hereby state that the content of the paper and computer readable forms of the Sequence Listing, submitted in the above-identified application in accordance with 37 C.F.R. § 1.821(c) and 1.821(e), respectively, are the same. The sequence listing is supported by the specification and references incorporated therein. Therefore, no new matter is added at this time.

Respectfully submitted,

Dated: Worden 11, 2002

30623 PATENT TRADEMARK OFFICE

c/o MINTZ, LEVIN, COHN, FERRIS, GLOVSKY and POPEO, P.C.

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Boston, Massachusetts 02111

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<211> 374

<212> PRT

<213> Homo sapiens

<400> 10

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Ser Glu Arg Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Leu Asn 20 25 30

| Pro | Ala | Arg 35 | Lys | Tyr | Phe | Pro | Gln 40 | Trp | Glu | Ala | Pro | Thr 45 | Leu | Val | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu | Gln 50 | Pro | Leu | Glu | Asn | Tyr 55 | Leu | Asp | Met | Glu | Tyr 60 | Phe | Gly | Thr | Ile |
| Gly 65 | Ile | Gly | Thr | Pro | Ala 70 | Gln | Asp | Phe | Thr | Val 75 | Val | Phe | Asp | Thr | Gly 80 |
| Ser | Ser | Asn | Leu | Trp 85 | Val | Pro | Ser | Val | Tyr 90 | Cys | Ser | Ser | Leu | Ala 95 | Cys |
| Thr | Asn | His | Asn 100 | Arg | Phe | Asn | Pro | Glu 105 | Asp | Ser | Ser | Thr | Tyr 110 | Gln | Ser |
| Thr | Ser | Glu 115 | Thr | Val | Ser | Ile | Thr 120 | Tyr | Gly | Thr | Gly | Ser 125 | Met | Thr | Gly |
| Ile | Leu 130 | Gly | Tyr | Asp | Thr | Val 135 | Gln | Val | Gly | Gly | Ile 140 | Ser | Asp | Thr | Asn |
| Gln 145 | Ile | Phe | Gly | Leu | Ser 150 | Glu | Thr | Glu | Pro | Gly 155 | Ser | Phe | Leu | Tyr | Tyr 160 |
| Ala | Pro | Phe | Asp | Gly 165 | Ile | Leu | Gly | Leu | Ala 170 | Tyr | Pro | Ser | Ile | Ser 175 | Ser |
| Ser | Gly | Ala | Thr 180 | Pro | Val | Phe | Asp | Asn 185 | Ile | Trp | Asn | Gln | Gly 190 | Leu | Val |
| Ser | Gln | Asp 195 | Leu | Phe | Ser | Val | Tyr 200 | Leu | Ser | Ala | Asp | Asp 205 | Lys | Ser | Gly |
| Ser | Val 210 | Val | Ile | Phe | Gly | Gly 215 | Ile | Asp | Ser | Ser | Tyr 220 | Tyr | Thr | Gly | Ser |
| Leu 225 | Asn | Trp | Val | Pro | Val 230 | Thr | Val | Glu | Gly | Tyr 235 | Trp | Gln | Ile | Thr | Val 240 |
| Asp | Ser | Ile | Thr | Met 245 | Asn | Gly | Glu | Thr | Ile 250 | Ala | Cys | Ala | Glu | Gly 255 | Cys |
| Gln | Ala | Ile | Val 260 | Asp | Thr | Gly | Thr | Ser 265 | Leu | Leu | Thr | Gly | Pro 270 | Thr | Ser |
| Pro | Ile | Ala 275 | Asn | Ile | Gln | Ser | Asp 280 | Ile | Gly | Ala | Ser | Glu 285 | Asn | Ser | Asp |

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Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile
290 295 300
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Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr 305 310 315 320

Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn 325 330 335

Val Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile 340 345 350

Arg Gln Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn Gln Val Ser Leu 355 360 365

Ala Pro Val Ala Val Asp 370

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<211> 1131

<212> DNA

<213> Homo sapiens

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<211> 374

<212> PRT

<213> Homo sapiens

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Pro Ala Arg Lys Tyr Phe Pro Gln Trp Glu Ala Pro Thr Leu Val Asp 35 40 45

Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile 50 55 60

Gly Ile Gly Thr Pro Ala Gln Asp Phe Thr Val Val Phe Asp Thr Gly 65 70 75 80

Ser Ser Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu Ala Cys 85 90 95

Thr Asn His Asn Arg Phe Asn Pro Glu Asp Ser Ser Thr Tyr Gln Ser 100 105 110

Thr Ser Glu Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met Thr Gly
115 120 125

Ile Leu Gly Tyr Asp Thr Val Gln Val Gly Gly Ile Ser Asp Thr Asn 130 135 140

Ala Pro Phe Asp Asp Ile Leu Gly Leu Ala Tyr Pro Ser Ile Ser Ser 165 170 175

Ser Gly Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly Leu Val 180 185 190

Ser Gln Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln Ser Gly
195 200 205

Ser Val Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr Gly Ser 210 215 220

Leu Asn Trp Val Pro Val Thr Val Glu Gly Tyr Trp Gln Ile Thr Val 225 230 235 240

Asp Ser Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu Gly Cys 245 250 255

Gln Ala Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Gly Pro Thr Ser 260 265 270

Pro Ile Ala Asn Ile Gln Ser Asp Ile Gly Ala Ser Glu Asn Ser Asp 275 280 285

Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile 290 295 300

Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr 305 310 315 320

Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn 325 330 335

Leu Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile 340 345 350

Arg Gln Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn Gln Val Ser Leu 355 360 365

Ala Pro Val Ala Val Asp 370

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<211> 479

<212> DNA

<213> Homo sapiens

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<211> 141

<212> PRT

<213> Homo sapiens

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<211> 326

<212> PRT

<213> Homo sapiens

<400> 16

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Ile Val His Leu Cys Leu Arg Lys Ala Asp Gln Lys Leu Val Ile Ile
20 25 30

Lys Gln Ile Pro Val Glu Gln Met Thr Lys Glu Glu Arg Gln Ala Ala 35 40 45

Gln Asn Glu Cys Gln Val Leu Lys Leu Leu Asn His Pro Asn Val Ile 50 55 60

Glu Tyr Tyr Glu Asn Phe Leu Glu Asp Lys Ala Leu Met Thr Ala Met 65 70 75 80

Glu Tyr Ala Pro Gly Gly Thr Leu Ala Glu Phe Ile Gln Lys Arg Cys 85 90 95

Asn Ser Leu Leu Glu Glu Glu Thr Ile Leu His Phe Phe Val Gln Ile 100 105 110

Leu Leu Ala Leu His His Val His Thr His Leu Ile Leu His Arg Asp 115 120 125

Leu Lys Thr Gln Asn Ile Leu Leu Asp Lys His Arg Met Val Val Lys
130 135 140

Ile Gly Asp Phe Gly Ile Ser Lys Ile Leu Ser Ser Lys Ser Lys Ala 145 150 155 160

Tyr Thr Val Val Gly Thr Pro Cys Tyr Ile Ser Pro Glu Leu Cys Glu 165 170 175 Gly Lys Pro Tyr Asn Gln Lys Ser Asp Ile Trp Ala Leu Gly Cys Val Leu Tyr Glu Leu Ala Ser Leu Lys Arg Ala Phe Glu Ala Ala Asn Leu Pro Ala Leu Val Leu Lys Ile Met Ser Gly Thr Phe Ala Pro Ile Ser Asp Arg Tyr Ser Pro Glu Leu Arg Gln Leu Val Leu Ser Leu Leu Ser Leu Glu Pro Ala Gln Arg Pro Pro Leu Ser His Ile Met Ala Gln Pro Leu Cys Ile Arg Ala Leu Leu Asn Leu His Thr Asp Val Gly Ser Val Arg Met Arg Arg Pro Val Gln Gly Gln Arg Ala Val Leu Gly Gly Arg Val Trp Ala Pro Ser Gly Ser Thr Leu Ser Pro Leu Thr Val Ser Ala Thr Ala Cys Thr Tyr Thr Leu Ser Ser Phe Thr Ile Asp Thr Leu His His Asp Leu Lys Thr Gln <210> 17

<400> 17

<211> 1591 <212> DNA

<213> Homo sapiens

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cagcacatca tccacagaga tgtcaagcct gacaacattc tcctggatga gagaggacat 720 gcacacctga ccgacttcaa cattgccacc atcatcaagg acggggagcg ggcgacggca 780 ttagcaggca ccaagccgta catggctccg gagatcttcc actcttttqt caacqqcqqq 840 accggctact ccttcgaggt ggactggtgg tcggtggggg tgatggccta tgagctgctg 900 cgaggatgga ggccctatga catccactcc agcaacqccq tgqagtccct ggtgcagctg 960 ttcagcaccg tgagcgtcca gtatgtcccc acgtggtcca aggagatggt qqqcttqctq 1020 cggaaggtgc tecteactgt gaaccecgag caecggetet ceagecteca qqaeqtqeag 1080 gcagcccgg cgctggccgg cgtgctgtgg gaccacctga gcgagaagag ggtggagccg 1140 ggcttcgtgc ccaacaaagg ccgtctgcac tgcqacccca cctttqaqct gqaqqaqatg 1200 atcctggagt ccaggcccct gcacaagaag aagaagcgcc tggccaagaa caagtcccgg 1260 gacaacagca gggacagete ecaqteegaq aatqactate tteaaqaetq eeteqatqee 1320 atccagcaag acttcgtgat ttttaacaga gaaaagctga agaggagcca ggacctcccg 1380 agggagcete teecegeece tgagtecagg gatgetgegg ageetgtgga ggacgaggeg 1440 gaacgctccg ccctgcccat gtgcggccc atttgcccct cggccgggag cggctaggcc 1500 gggacgcccg tggtcctcac cccttgagct gctttggaga ctcggctgcc agagggaggg 1560 ccatgggccg aggcctggca ttcacgttcc c 1591

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<211> 488

<212> PRT

<213> Homo sapiens

<400> 18

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Ser Ala Leu Pro Pro Pro Ala Ala Gly Gln Pro Arg Ala Arg Asp Ser 35 40 45

Gly Asp Val Arg Ser Gln Pro Arg Pro Leu Phe Gln Trp Ser Lys Trp 50 55 60

Lys Lys Arg Met Gly Ser Ser Met Ser Ala Ala Thr Ala Arg Arg Pro 65 70 75 80

Val Phe Asp Asp Lys Glu Asp Val Asn Phe Asp His Phe Gln Ile Leu 85 90 95

Arg Ala Ile Gly Lys Gly Ser Phe Gly Lys Val Val Cys Ile Val Gln 100 105 110

Lys Arg Asp Thr Glu Lys Met Tyr Ala Met Lys Tyr Met Asn Lys Gln
115 120 125

| Gln | Cys 130 | Ile | Glu | Arg | Asp | Glu 135 | Val | Arg | Asn | Val | Phe 140 | Arg | Glu | Leu | Glu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile 145 | Leu | Gln | Glu | Ile | Glu 150 | His | Val | Phe | Leu | Val 155 | Asn | Leu | Trp | Tyr | Ser 160 |
| Phe | Gln | Asp | Glu | Glu 165 | Asp | Met | Phe | Met | Val 170 | Val | Asp | Leu | Leu | Leu 175 | Gly |
| Gly | Asp | Leu | Arg 180 | Tyr | His | Leu | Gln | Gln 185 | Asn | Val | Gln | Phe | Ser 190 | Glu | Asp |
| Thr | Val | Arg 195 | Leu | Tyr | Ile | Cys | Glu 200 | Met | Ala | Leu | Ala | Leu 205 | Asp | Tyr | Leu |
| Arg | Gly 210 | Gln | His | Ile | Ile | His 215 | Arg | Asp | Val | Lys | Pro 220 | Asp | Asn | Ile | Leu |
| Leu 225 | Asp | Glu | Arg | Gly | His 230 | Ala | His | Leu | Thr | Asp 235 | Phe | Asn | Ile | Ala | Thr 240 |
| Ile | Ile | Lys | Asp | Gly 245 | Glu | Arg | Ala | Thr | Ala 250 | Leu | Ala | Gly | Thr | Lys 255 | Pro |
| Tyr | Met | Ala | Pro 260 | Glu | Ile | Phe | His | Ser 265 | Phe | Val | Asn | Gly | Gly 270 | Thr | Gly |
| Tyr | Ser | Phe 275 | Glu | Val | Asp | Trp | Trp 280 | Ser | Val | Gly | Val | Met 285 | Ala | Tyr | Glu |
| Leu | Leu 290 | Arg | Gly | Trp | Arg | Pro 295 | Tyr | Asp | Ile | His | Ser 300 | Ser | Asn | Ala | Val |
| Glu 305 | Ser | Leu | Val | Gln | Leu 310 | Phe | Ser | Thr | Val | Ser 315 | Val | Gln | Tyr | Val | Pro 320 |
| Thr | Trp | Ser | Lys | Glu 325 | Met | Val | Gly | Leu | Leu 330 | Arg | Lys | Val | Leu | Leu 335 | Thr |
| Val | Asn | Pro | Glu 340 | His | Arg | Leu | Ser | Ser 345 | Leu | Gln | Asp | Val | Gln 350 | Ala | Ala |
| Pro | Ala | Leu 355 | Ala | Gly | Val | Leu | Trp 360 | Asp | His | Leu | Ser | Glu 365 | Lys | Arg | Val |
| Glu | Pro 370 | Gly | Phe | Val | Pro | Asn 375 | Lys | Gly | Arg | Leu | His 380 | Cys | Asp | Pro | Thr |

Phe Glu Leu Glu Glu Met Ile Leu Glu Ser Arg Pro Leu His Lys Lys 385 390 395 400 Lys Lys Arg Leu Ala Lys Asn Lys Ser Arg Asp Asn Ser Arg Asp Ser 405 410 415 Ser Gln Ser Glu Asn Asp Tyr Leu Gln Asp Cys Leu Asp Ala Ile Gln 425 420 430 Gln Asp Phe Val Ile Phe Asn Arg Glu Lys Leu Lys Arg Ser Gln Asp 435 440 445 Leu Pro Arg Glu Pro Leu Pro Ala Pro Glu Ser Arg Asp Ala Ala Glu 450 455 460 Pro Val Glu Asp Glu Ala Glu Arg Ser Ala Leu Pro Met Cys Gly Pro 470 475 465 480 Ile Cys Pro Ser Ala Gly Ser Gly 485 <210> 19 <211> 581 <212> DNA <213> Homo sapiens <400> 19 cactccagag ctcagagcca cccacagcca cagctatgca gtgcctcctg ctcaccctga 60 qcatqqccct qqtctqtqcc atccagqcca gggacatccc ccagaccaag caggacqtgg 120 ageteceaaa gttggeaggg acetggtact ceatggeeat ggtggeeagt gaettetece 180 tectggagae egtggaggee cetetgaggg teaacateae etegetgtgg eccaeeceeg 240 agggcaacct ggagatcatt ctgcacagat gggaacacca cagatgcgtt gagaggaccg 300 tectequeca quaquetqua queceggetg tgtteatggt egacegtage aggagetaeg 360 tqttcttctq catqqqqacc accacacca gtqctqacca ccacacqatg tqccagtacc 420 tqqqqatqac aqccaqqacc ctagaqqcag acqacaaqqt catqqaqqaa ttcatcaqct 480 ttctcaggac cctgcccgtg cacatgtgga tcttcctgga cgttacccag gcggaacagt 540 gccqcqtcta qatqaqctcc tqctcaqtcc tgcctcctqq q 581 <210> 20 <211> 171 <212> PRT <213> Homo sapiens <400> 20 Met Gln Cys Leu Leu Thr Leu Ser Met Ala Leu Val Cys Ala Ile

10

15

5

Gln Ala Arg Asp Ile Pro Gln Thr Lys Gln Asp Val Glu Leu Pro Lys 20 25 Leu Ala Gly Thr Trp Tyr Ser Met Ala Met Val Ala Ser Asp Phe Ser 35 40 45 Leu Leu Glu Thr Val Glu Ala Pro Leu Arg Val Asn Ile Thr Ser Leu 55 60 Trp Pro Thr Pro Glu Gly Asn Leu Glu Ile Ile Leu His Arg Trp Glu 65 70 75 His His Arg Cys Val Glu Arg Thr Val Leu Ala Gln Lys Thr Glu Asp 85 90 Pro Ala Val Phe Met Val Asp Arg Ser Arg Ser Tyr Val Phe Cys 100 105 Met Gly Thr Thr Pro Ser Ala Asp His His Thr Met Cys Gln Tyr 115 120 125 Leu Gly Met Thr Ala Arg Thr Leu Glu Ala Asp Asp Lys Val Met Glu 135 Glu Phe Ile Ser Phe Leu Arg Thr Leu Pro Val His Met Trp Ile Phe 150 155 160 Leu Asp Val Thr Gln Ala Glu Gln Cys Arg Val 165 170

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<213> Homo sapiens

<400> 21

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Phe Arg Arg Leu Arg Gln Phe Arg Gln Ala Gln Pro Thr Pro Gln Tyr 35 40 45

Arg Phe Arg Lys Arg Asp Lys Val Met Phe Tyr Gly Arg Lys Ile Met 50 55 60

Arg Lys Val Thr Thr Leu Pro Asn Thr Leu Val Glu Asn Thr Ala Leu 65 70 75 80

Pro Arg Gln Arg Ala Arg Lys Arg Thr Lys Val Leu Ser Leu Ala Lys 85 90 95

Arg Ile Leu Arg Phe Lys Lys Glu Tyr Pro Ala Leu Gln Pro Lys Glu

| | | | 100 | | | | | 105 | | | | | 110 | | |
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| Pro | Pro | Pro 115 | Ser | Leu | Leu | Glu | Ala 120 | Asp | Leu | Thr | Glu | Phe 125 | Asp | Val | Lys |
| Asn | Ser 130 | His | Leu | Pro | Ser | Glu 135 | Val | Leu | Tyr | Met | Leu 140 | Lys | Asn | Val | Arg |
| Val 145 | Leu | Gly | His | Phe | Glu 150 | Lys | Pro | Leu | Phe | Leu 155 | Glu | Leu | Cys | Lys | His 160 |
| Ile | Val | Phe | Val | Gln 165 | Leu | Gln | Glu | Gly | Glu 170 | His | Val | Phe | Gln | Pro 175 | Arg |
| Glu | Pro | Asp | Pro 180 | Ser | Ile | Cys | Val | Val 185 | Gln | Asp | Gly | Arg | Leu 190 | Glu | Val |
| Cys | Ile | Gln 195 | Asp | Thr | Asp | Gly | Thr 200 | Glu | Val | Val | Val | Lys 205 | Glu | Val | Leu |
| Ala | Gly 210 | Asp | Ser | Val | His | Ser 215 | Leu | Leu | Ser | Ile | Leu 220 | Asp | Ile | Ile | Thr |
| Gly 225 | His | Ala | Ala | Pro | Tyr 230 | Lys | Thr | Val | Ser | Val 235 | Arg | Ala | Ala | Ile | Pro 240 |
| Ser | Thr | Ile | Leu | Arg 245 | Leu | Pro | Ala | Ala | Ala 250 | Phe | His | Gly | Val | Phe 255 | Glu |
| Lys | Tyr | Pro | Glu 260 | Thr | Leu | Val | Arg | Val 265 | Val | Gln | Leu | Gln | Ile 270 | Ile | Met |
| Val | Arg | Leu 275 | Gln | Arg | Val | Thr | Phe 280 | Leu | Ala | Leu | His | Asn 285 | Tyr | Leu | Gly |
| Leu | Thr 290 | Thr | Glu | Leu | Phe | Asn 295 | Ala | Glu | Ser | Gln | Ala 300 | Ile | Pro | Leu | Val |
| Ser 305 | Val | Ala | Ser | Val | Ala 310 | Ala | Gly | Lys | Ala | Lys 315 | Lys | Gln | Val | Phe | Tyr 320 |
| Gly | Glu | Glu | Glu | Arg 325 | Leu | Lys | Lys | Pro | Pro 330 | Arg | Leu | His | Glu | Ser 335 | Cys |
| Asp | Ser | Ala | Asp 340 | His | Gly | Gly | Gly | Arg 345 | Pro | Ala | Ala | Ala | Gly 350 | Pro | Leu |

Leu Lys Arg Ser His Ser Val Pro Ala Pro Ser Ile Arg Lys Gln Ile

| T.e.11 | Glu | Glu | Leu | Glu | Lvs | Pro | Glv | Ala | Glv | Asp | Pro | Asp | Pro | |
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360

355

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- Pro Gln Gly Gly Pro Gly Ser Ala Thr Ser Asp Leu Gly Met Ala Cys 385 390 395 400
- Asp Arg Ala Arg Val Phe Leu His Ser Asp Glu His Pro Gly Ser Ser 405 410 415
- Val Ala Ser Lys Ser Arg Lys Ser Val Met Val Ala Glu Ile Pro Ser 420 425 430
- Thr Val Ser Gln His Ser Glu Ser His Thr Asp Glu Thr Leu Ala Ser 435 440 445
- Arg Lys Ser Asp Ala Ile Phe Arg Ala Ala Lys Lys Asp Leu Leu Thr 450 455 460
- Leu Met Lys Leu Glu Asp Ser Ser Leu Leu Asp Gly Arg Val Ala Leu 465 470 475 480
- Leu His Val Pro Ala Cys Thr Val Val Ser Met Gln Gly Asp Gln Asp
 485
 490
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- Ala Ser Ile Leu Phe Val Val Leu Gly Leu Leu His Val Tyr Gln Arg
 500 505 510
- Lys Ile Cys Ser Gln Glu Asp Thr Cys Leu Phe Ser Arg Ala Pro Gly 515 520 525
- Asp Ser Ser Leu Leu Asp Gly Arg Val Ala Leu Leu His Val Pro Ala 530 535 540
- Gly Thr Val Val Ser Arg Gln Gly Asp Gln Asp Ala Ser Ile Leu Phe 545 550 555 560
- Val Val Ser Gly Leu Leu His Val Tyr Gln Arg Lys Ile Gly Ser Gln 565 570 575
- Glu Asp Thr Cys Leu Phe Leu Thr Arg Pro Gly Glu Met Val Gly Gln
 580 585 590
- Leu Ala Val Leu Thr Gly Glu Pro Leu Ile Phe Thr Val Lys Ala Asn 595 600 605
- Arg Asp Cys Ser Phe Leu Ser Ile Ser Lys Ala His Phe Tyr Glu Ile

| 610 | 615 | 620 |
|-----|-----|-----|
| | | |

| Met 625 | Arg | Lys | Gln | Pro | Thr 630 | Val | Val | Leu | Gly | Val 635 | Ala | His | Thr | Val | Val 640 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Lys | Arg | Met | Ser | Ser 645 | Phe | Val | Arg | Gln | Ile 650 | Asp | Phe | Ala | Leu | Asp 655 | Trp |
| Val | Glu | Val | Glu 660 | Ala | Gly | Arg | Ala | Ile 665 | Tyr | Arg | Gln | Gly | Asp 670 | Lys | Ser |
| Asp | Cys | Thr 675 | Tyr | Ile | Met | Leu | Ser 680 | Gly | Arg | Leu | Arg | Ser 685 | Val | Ile | Arg |
| Lys | Asp 690 | Asp | Gly | Lys | Lys | Arg 695 | Leu | Ala | Gly | Glu | Tyr 700 | Gly | Arg | Gly | Asp |
| Leu 705 | Val | Gly | Val | Val | Glu 710 | Thr | Leu | Thr | His | Gln 715 | Ala | Arg | Ala | Thr | Thr 720 |
| Val | His | Ala | Val | Arg 725 | Asp | Ser | Glu | Leu | Ala 730 | Lys | Leu | Pro | Ala | Gly 735 | Ala |
| Leu | Thr | Cys | Ile 740 | Lys | Arg | Arg | Tyr | Pro 745 | Gln | Val | Val | Thr | Arg 750 | Leu | Ile |
| His | Leu | Leu 755 | Gly | Glu | Lys | Ile | Leu 760 | Gly | Ser | Leu | Gln | Gln 765 | Gly | Pro | Val |
| Thr | Gly 770 | His | Gln | Leu | Gly | Leu 775 | Pro | Thr | Glu | Gly | Ser 780 | Lys | Trp | Asp | Leu |
| Gly 785 | Asn | Pro | Ala | Val | Asn 790 | Leu | Ser | Thr | Val | Ala 795 | Val | Met | Pro | Val | Ser 800 |
| Glu | Glu | Val | Pro | Leu 805 | Thr | Ala | Phe | Ala | Leu 810 | Glu | Leu | Glu | His | Ala 815 | Leu |
| Ser | Ala | Ile | Gly 820 | Pro | Pro | Leu | Leu | Leu 825 | Thr | Ser | Asp | Asn | Ile 830 | Lys | Arg |
| Arg | Leu | Gly 835 | Ser | Ala | Ala | Leu | Asp 840 | Ser | Val | His | Glu | Tyr 845 | Arg | Leu | Ser |
| Ser | Trp 850 | Leu | Gly | Gln | Gln | Glu 855 | Asp | Thr | His | Arg | Ile 860 | Val | Leu | Tyr | Gln |

Val Asp Gly Thr Leu Thr Pro Trp Thr Gln Arg Cys Val Arg Gln Ala

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- Glu Leu Glu Arg Met Leu Glu Ser Thr Ala Val Arg Ala Gln Lys Gln 900 905 910
- Leu Ile Leu Leu His Arg Glu Glu Gly Pro Ala Pro Ala Arg Thr Val 915 920 925
- Glu Trp Leu Asn Met Arg Ser Trp Cys Ser Gly His Leu His Leu Cys 930 935 940
- Cys Pro Arg Arg Val Phe Ser Arg Arg Ser Leu Pro Lys Leu Val Glu 945 950 955 960
- Met Tyr Lys His Val Phe Gln Arg Pro Pro Asp Arg His Ser Asp Phe 965 970 975
- Ser Arg Leu Ala Arg Val Leu Thr Gly Asn Ala Ile Ala Leu Val Leu 980 985 990
- Gly Gly Gly Ala Ser Met Thr Ser Leu Met Lys Ala Ala Leu Asp 995 1000 1005
- Leu Thr Tyr Pro Ile Thr Ser Met Phe Ser Gly Ala Gly Phe Asn Ser 1010 1015 1020
- Ser Ile Phe Ser Val Phe Lys Asp Gln Gln Ile Glu Asp Leu Trp Ile 1025 1030 1035 1040
- Pro Tyr Phe Ala Ile Thr Thr Asp Ile Thr Ala Ser Ala Met Arg Val $1045 \hspace{1.5cm} 1050 \hspace{1.5cm} 1055$
- His Thr Asp Gly Ser Leu Trp Trp Tyr Val Arg Ala Ser Met Ser Leu 1060 1065 1070
- Ser Gly Tyr Met Pro Pro Leu Cys Asp Pro Lys Asp Gly His Leu Leu 1075 1080 1085
- Met Asp Gly Gly Tyr Ile Asn Asn Leu Pro Ala Ala Ser Ala Pro Arg 1090 1095 1100
- Ser Leu Gly Trp Asn Thr Phe Ser Leu Glu Tyr Ala Lys Gly Lys Cys 1105 1110 1115 1120
- Gln Ala Gly Ile Arg Ala Pro Arg Thr Cys Thr Arg Val Tyr Met His

| 1125 1130 1135 |
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- Thr Gln Ala Pro Ala Ala Cys Ala Pro Ala Tyr Gly Pro Val Cys Gln 1140 1145 1150
- Leu Ser Ser Met Gln Asn Lys Gly Gln Val Glu Glu Leu Gly Ala Ile 1155 1160 1165
- Lys Pro His Leu Cys Pro Gln Ser Glu Thr Asn Ser Leu Gln Gly Val 1170 1175 1180
- Thr Arg Ala Gly Phe Ser Leu Ala Asp Val Ala Arg Ser Met Gly Ala 1185 1190 1195 1200
- Lys Val Val Ile Ala Ile Asp Val Gly Ser Arg Asp Glu Thr Asp Leu 1205 1210 1215
- Thr Asn Tyr Gly Asp Ala Leu Ser Gly Trp Trp Leu Leu Trp Lys Arg 1220 1225 1230
- Trp Asn Pro Leu Ala Thr Lys Val Lys Val Leu Asn Met Ala Glu Ile 1235 1240 1245
- Gln Thr Arg Leu Ala Tyr Val Cys Cys Val Arg Gln Leu Glu Val Val 1250 1255 1260
- Lys Ser Ser Asp Tyr Cys Glu Tyr Leu Arg Pro Pro Ile Asp Ser Tyr 1265 1270 1275 1280
- Ser Thr Leu Asp Phe Gly Lys Phe Asn Glu Ile Cys Glu Val Gly Tyr 1285 1290 1295
- Gln His Gly Arg Thr Val Phe Asp Ile Trp Gly Arg Ser Gly Val Leu 1300 1305 1310
- Glu Lys Met Leu Arg Asp Gln Gln Gly Pro Ser Lys Lys Pro Ala Ser 1315 1320 1325
- Ala Val Leu Thr Cys Pro Asn Ala Ser Phe Thr Asp Leu Ala Glu Ile 1330 1335 1340
- Val Ser Arg Ile Glu Pro Ala Lys Pro Ala Met Val Asp Asp Glu Ser 1345 1350 1355 1360
- Asp Tyr Gln Thr Glu Tyr Glu Glu Glu Leu Leu Asp Val Pro Arg Asp 1365 1370 1375
- Ala Tyr Ala Asp Phe Gln Ser Thr Ser Ala Gln Gln Gly Ser Asp Leu

1380 1385 1390

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Ala Glu Pro His Arg Ala Gly Arg Gln Trp Lys Phe Pro Gly Ser Phe 50 55 60

Tyr Phe Ala Ile Thr Val Ile Thr Thr Ile Glu Tyr Gly His Ala Ala 65 70 75 80 Pro Gly Thr Asp Ser Gly Lys Val Phe Cys Met Phe Tyr Ala Leu Leu 85 90 95 Gly Ile Pro Leu Thr Leu Val Thr Phe Gln Ser Leu Gly Glu Arg Leu 100 105 110 Asn Ala Val Val Arg Arg Leu Leu Leu Ala Ala Lys Cys Cys Leu Gly 115 120 125 Leu Arg Trp Thr Cys Val Ser Thr Glu Asn Leu Val Val Ala Gly Leu 130 135 140 Leu Ala Cys Ala Ala Thr Leu Ala Leu Gly Ala Val Ala Phe Ser His 145 150 155 Phe Glu Gly Trp Thr Phe Phe His Ala Tyr Tyr Tyr Cys Phe Ile Thr 170 165 Leu Thr Thr Ile Gly Phe Gly Asp Asn Leu Gly Phe Ser Pro Pro Ser 185 Ser Pro Gly Val Val Arg Gly Gln Ala Pro Arg Leu Gly Ala Arg 200 205 Trp Lys Ser Ile 210 <210> 25 <211> 711 <212> DNA <213> Homo sapiens <400> 25 tetetetete tetetetete tetggtgaac aggaceegte gecatgggee gtgtgateeg 60 tggacagagg aagggcgccg ggtctgtgtt ccgcgcgcac gtgaagcacc gtaaaggcgc 120 tgcgcgcctg cgcgccgtgg atttcgctga gcggcacggc tacatcaagg gcatcgtcaa 180

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|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Lys | Ala | Ala 35 | Gln | Pro | Tyr | Leu | Arg 40 | Arg | Gln | Arg | Leu | Leu 45 | Arg | Asp | Leu |
| Arg | Pro 50 | Phe | Pro | Ala | Pro | Pro 55 | Thr | His | Trp | Phe | Leu 60 | Gly | His | Lys | Leu |
| Met 65 | Glu | Lys | Tyr | Pro | Cys 70 | Ala | Val | Pro | Leu | Trp 75 | Val | Gly | Pro | Phe | Thr 80 |
| Met | Phe | Phe | Ser | Val 85 | His | Asp | Pro | Asp | Tyr 90 | Ala | Lys | Ile | Leu | Leu 95 | Lys |
| Arg | Gln | Gly | Lys 100 | Asn | Gln | Glu | Gly | Phe 105 | Leu | Pro | Phe | Ile | Ser 110 | Gln | Gly |
| Lys | Gly | Leu 115 | Ala | Ala | Leu | Asp | Gly 120 | Pro | Lys | Trp | Phe | Gln 125 | His | Arg | Arg |
| Leu | Leu 130 | Thr | Pro | Gly | Phe | His 135 | Phe | Asn | Ile | Leu | Lys 140 | Ala | Tyr | Ile | Glu |
| Val 145 | Met | Ala | His | Ser | Val 150 | Lys | Met | Met | Leu | Asn 155 | Lys | Trp | Glu | Glu | His 160 |
| Ile | Ala | Gln | Asn | Ser 165 | Arg | Leu | Glu | Leu | Phe 170 | Gln | His | Val | Ser | Leu 175 | Met |
| Thr | Leu | Asp | Ser 180 | Ile | Met | Lys | Cys | Ala 185 | Phe | Ser | His | Gln | Gly 190 | Ser | Ile |
| Gln | Leu | Asp 195 | Arg | Ser | Ser | Tyr | Leu 200 | Lys | Ala | Val | Phe | Asn 205 | Leu | Ser | Lys |
| Ile | Ser 210 | Asn | Gln | Arg | Met | Asn 215 | Asn | Phe | Leu | His | His 220 | Asn | Asp | Leu | Val |
| Phe 225 | Lys | Phe | Ser | Ser | Gln 230 | Gly | Gln | Ile | Phe | Ser 235 | Lys | Phe | Asn | Gln | Glu 240 |
| Leu | His | Gln | His | Leu 245 | Glu | Lys | Val | Ile | Gln 250 | Asp | Arg | Lys | Glu | Ser 255 | Leu |
| Lys | Asp | Lys | Leu 260 | Lys | Gln | Asp | Thr | Thr 265 | Gln | Lys | Arg | Arg | Trp 270 | Asp | Phe |

Leu Asp Ile Leu Leu Ser Ala Lys Val Glu Asn Thr Lys Asp Phe Ser 275 280 285

Glu Ala Asp Leu Gln Ala Glu Val Lys Thr Phe Met Phe Ala Gly His 290 295 300

Asp Thr Thr Ser Ser Ala Ile Ser Trp Ile Leu Tyr Cys Leu Ala Lys 305 310 315 320

Tyr Pro Glu His Gln Gln Arg Cys Arg Asp Glu Ile Arg Glu Leu Leu 325 · 330 335

Gly Asp Gly Ser Ser Ile Thr Trp His Leu Ser Gln Met Pro Tyr Thr 340 345 350

Thr Met Cys Ile Lys Glu Cys Leu Arg Leu Tyr Ala Pro Val Val Asn 355 360 365

Ile Ser Arg Leu Leu Asp Lys Pro Ile Thr Phe Pro Asp Gly Arg Ser 370 380

Leu Pro Ala Gly Ile Thr Val Val Leu Ser Ile Trp Gly Leu His His 385 390 395 400

Asn Pro Ala Val Trp Lys Asn Val Gln Val Phe Asp Pro Leu Arg Phe 405 410 415

Ser Gln Glu Asn Ser Asp Gln Arg His Pro Tyr Ala Tyr Leu Pro Phe 420 425 430

Ser Ala Gly Ser Arg Asn Cys Ile Gly Gln Glu Phe Ala Met Ile Glu 435 440 445

Leu Lys Val Thr Ile Ala Leu Ile Leu Leu His Phe Arg Val Thr Pro 450 455 460

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Gly Cys Thr Thr Phe Ser Leu Val Ala His Arg Gly Gly Phe Ala Gly
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                             40
Val Gln Gly Thr Phe Cys Met Asp Ala Trp Gly Phe Cys Phe Ala Val
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Ser Ala Leu Val Val Ala Cys Glu Phe Thr Arg Leu His Gly Cys Leu
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| Arg | Leu | Ser | Trp | Gly 85 | Asn | Phe | Thr | Ala | Ala 90 | Phe | Ala | Met | Leu | Ala 95 | Thr |
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| Leu | Leu | Cys | Ala 100 | Thr | Ala | Ala | Val | Leu 105 | Tyr | Pro | Leu | Tyr | Phe 110 | Ala | Arg |
| Arg | Glu | Cys 115 | Ser | Pro | Glu | Pro | Ala 120 | Gly | Cys | Ala | Ala | Arg 125 | Asp | Phe | Arg |
| Leu | Ala 130 | Ala | Ser | Val | Phe | Ala 135 | Gly | Leu | Leu | Phe | Leu 140 | Ala | Tyr | Ala | Val |
| Glu 145 | Val | Ala | Leu | Thr | Arg 150 | Ala | Arg | Pro | Gly | Gln 155 | Val | Ser | Ser | Tyr | Met 160 |
| Ala | Thr | Val | Ser | Gly 165 | Leu | Leu | Lys | Ile | Val 170 | Gln | Ala | Phe | Val | Ala 175 | Суѕ |
| Ile | Ile | Phe | Gly 180 | Ala | Leu | Val | His | Asp 185 | Ser | Arg | Tyr | Gly | Arg 190 | Tyr | Val |
| Ala | Thr | Gln 195 | Trp | Cys | Val | Ala | Val 200 | Tyr | Ser | Leu | Cys | Phe 205 | Leu | Ala | Thr |
| Val | Ala 210 | Val | Val | Ala | Leu | Ser 215 | Val | Met | Gly | His | Thr 220 | Gly | Gly | Leu | Gly |
| Cys 225 | Pro | Phe | Asp | Arg | Leu 230 | Val | Val | Val | Tyr | Thr 235 | Phe | Leu | Ala | Val | Leu 240 |
| Leu | Tyr | Leu | Ser | Ala 245 | Ala | Val | Ile | Trp | Pro 250 | Val | Phe | Cys | Phe | Asp 255 | Pro |
| Lys | Tyr | Gly | Glu 260 | Pro | Lys | Arg | Pro | Pro 265 | Asn | Cys | Ala | Arg | Gly 270 | Ser | Cys |
| Pro | Trp | Asp 275 | Thr | Ser | Trp | Trp | Trp 280 | Pro | Ser | Ser | Pro | Thr 285 | Ser | Thr | Cys |
| Ser | Cys 290 | Thr | Ser | Leu | Thr | Ser 295 | Pro | Thr | Pro | Ser | Phe 300 | Ser | Ser | Ala | Arg |
| Arg 305 | Ala | Ser | Val | His | Cys 310 | Gly | His | Leu | Trp | His 315 | Trp | Glu | Gly | Ala | Arg 320 |
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55

50

Leu Met Phe Pro Lys Glu Lys Glu Ala Phe Leu Ala Leu Ala Gln Leu

60

| Leu 65 | Thr | Ser | Lys | Asn | Leu 70 | Pro | Asp | Thr | Val | Asp 75 | Gly | Gln | Leu | Pro | Met 80 |
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| Gly | Pro | His | Ser | Arg 85 | Ala | Ser | Gln | Val | Ala 90 | Pro | Glu | Thr | Thr | Ser 95 | Ser |
| Lys | Val | Asp | Arg 100 | Gly | Val | Ser | Thr | Val 105 | Cys | Gly | Lys | Pro | Lys 110 | Val | Val |
| Gly | Lys | Ile 115 | Tyr | Gly | Gly | Arg | Asp 120 | Ala | Ala | Ala | Gly | Gln 125 | Trp | Pro | Trp |
| Gln | Ala 130 | Ser | Leu | Leu | Tyr | Trp 135 | Gly | Ser | His | Leu | Cys 140 | Gly | Ala | Val | Leu |
| Ile 145 | Asp | Ser | Cys | Trp | Leu 150 | Val | Ser | Thr | Thr | His 155 | Cys | Phe | Lys | Ser | Gln 160 |
| Ala | Pro | Lys | Asn | Tyr 165 | Gln | Val | Leu | Leu | Gly 170 | Asn | Ile | Gln | Leu | Tyr 175 | His |
| Gln | Thr | Gln | His 180 | Thr | Gln | Lys | Met | Ser 185 | Val | His | Arg | Ile | Ile 190 | Thr | His |
| Pro | Asp | Phe 195 | Glu | Lys | Leu | His | Pro 200 | Phe | Gly | Ser | Asp | Ile 205 | Ala | Met | Leu |
| Gln | Leu 210 | His | Leu | Pro | Met | Asn 215 | Phe | Thr | Ser | Tyr | Ile 220 | Val | Pro | Val | Cys |
| Leu 225 | Pro | Ser | Arg | Asp | Met 230 | Gln | Leu | Pro | Ser | Asn 235 | Val | Ser | Cys | Trp | Ile 240 |
| Thr | Gly | Trp | Gly | Met 245 | Leu | Thr | Glu | Asp | Leu 250 | Cys | Ser | Gln | Gly | Asp 255 | Ser |
| Gly | Gly | Pro | Leu 260 | Val | Cys | Tyr | Leu | Pro 265 | Ser | Ala | Trp | Val | Leu 270 | Val | Gly |
| Leu | Ala | Ser 275 | Trp | Gly | Leu | Asp | Cys 280 | Arg | His | Pro | Ala | Tyr 285 | Pro | Ser | Ile |
| Phe | Thr 290 | Arg | Val | Thr | Tyr | Phe 295 | Ile | Asn | Trp | Ile | Asp 300 | Lys | Ile | Met | Arg |
| Leu 305 | Thr | Pro | Leu | Ser | Asp 310 | Pro | Ala | Leu | Ala | Pro 315 | His | Thr | Cys | Ser | Pro 320 |

Pro Lys Pro Leu Arg Ala Ala Gly Leu Pro Gly Pro Cys Ala Ala Leu 325 330 335

Val Leu Pro Gln Thr Trp Leu Leu Pro Leu Thr Leu Arg Ala Pro 340 345 350

Trp Gln Thr Leu 355

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Trp Leu Ala Glu Gln Arg Val Pro Leu Ser Val Gln Leu Lys Pro Glu 35 40 45

Val Ser Pro Thr Gln Asp Ile Arg Leu Trp Val Ser Val Glu Asp Ala
50 55 60

Gln Met His Thr Val Thr Ile Trp Leu Thr Val Arg Pro Asp Met Thr
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Val Ala Ser Leu Lys Asp Met Val Phe Leu Asp Tyr Gly Phe Pro Pro 85 90 95

Val Leu Gln Gln Trp Val Ile Gly Gln Arg Leu Ala Arg Asp Gln Glu 100 105 110

Thr Leu His Ser His Gly Val Arg Gln Asn Gly Asp Ser Ala Tyr Leu 115 120 125

Tyr Leu Leu Ser Ala Arg Asn Thr Ser Leu Asn Pro Gln Glu Leu Gln 130 135 140

Arg Glu Arg Gln Leu Arg Met Leu Glu Asp Leu Gly Phe Lys Asp Leu 145 150 155 160

Thr Leu Gln Pro Arg Gly Pro Leu Glu Pro Gly Pro Pro Lys Pro Gly

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| Pro | Pro | Val 195 | Gly | Trp | Gln | Cys | Pro 200 | Gly | Cys | Thr | Phe | Ile 205 | Asn | Lys | Pro |
| Thr | Arg 210 | Pro | Gly | Cys | Glu | Met 215 | Cys | Cys | Arg | Ala | Arg 220 | Pro | Glu | Ala | Tyr |
| Gln 225 | Val | Pro | Ala | Ser | Tyr 230 | Gln | Pro | Asp | Glu | Glu 235 | Glu | Arg | Ala | Arg | Leu 240 |
| Ala | Gly | Glu | Glu | Glu 245 | Ala | Leu | Arg | Gln | Tyr 250 | Gln | Gln | Arg | Lys | Gln 255 | Gln |
| Gln | Gln | Glu | Gly 260 | Asn | Tyr | Leu | Gln | His 265 | Val | Gln | Leu | Asp | Gln 270 | Arg | Ser |
| Leu | Val | Leu 275 | Asn | Thr | Glu | Pro | Ala 280 | Glu | Cys | Pro | Val | Cys 285 | Tyr | Ser | Val |
| Leu | Ala 290 | Pro | Gly | Glu | Ala | Val 295 | Val | Leu | Arg | Glu | Cys 300 | Leu | His | Thr | Phe |
| Cys 305 | Arg | Glu | Cys | Leu | Gln 310 | Gly | Thr | Ile | Arg | Asn 315 | Ser | Gln | Glu | Ala | Glu 320 |
| Val | Ser | Cys | Pro | Phe 325 | Ile | Asp | Asn | Thr | Tyr 330 | Ser | Cys | Ser | Gly | Lys 335 | Leu |
| Leu | Glu | Arg | Glu 340 | Ile | Lys | Ala | Leu | Leu 345 | Thr | Pro | Glu | Asp | Tyr 350 | Gln | Arg |
| Phe | Leu | Asp 355 | Leu | Gly | Ile | Ser | Ile 360 | Ala | Glu | Asn | Arg | Ser 365 | Ala | Phe | Ser |
| Tyr | His 370 | Cys | Lys | Thr | Pro | Asp 375 | Cys | Lys | Gly | Trp | Cys 380 | Phe | Phe | Glu | Asp |
| Asp 385 | Val | Asn | Glu | Phe | Thr 390 | Cys | Pro | Val | Cys | Phe 395 | His | Val | Asn | Cys | Leu 400 |
| Leu | Cys | Lys | Ala | Ile 405 | His | Glu | Gln | Met | Asn 410 | Cys | Lys | Glu | Tyr | Gln 415 | Glu |
| _ | _ | | _ | _ | | | _ | _ | | | | _ | | | |

Asp Leu Ala Leu Arg Ala Gln Asn Asp Val Ala Ala Arg Gln Thr Thr

420 425 430

Glu Met Leu Lys Val Met Leu Gln Gln Gly Glu Ala Met Arg Cys Pro $435 \hspace{1.5cm} 440 \hspace{1.5cm} 445 \hspace{1.5cm}$

Gln Cys Gln Ile Val Val Gln Lys Lys Asp Gly Cys Asp Trp Ile Arg 450 460

Cys Thr Val Cys His Thr Glu Ile Cys Trp Val Thr Lys Gly Pro Arg 465 470 475 480

Trp Gly Pro Gly Pro Gly Asp Thr Ser Gly Gly Cys Arg Cys Arg
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Val Asn Gly Ile Pro Cys His Pro Ser Cys Gln Asn Cys His
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<211> 2372

<212> DNA

<213> Homo sapiens

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<212> PRT

<213> Homo sapiens

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Ala Arg Ser Pro Pro Gly Gly Trp Ala Gln Pro Arg Gln Met Asp Glu
35 40 45

Lys Thr Lys Lys Ala Glu Glu Met Ala Leu Ser Leu Thr Arg Ala Val 50 55 60

Ala Gly Gly Asp Glu Gln Val Ala Met Lys Cys Ala Ile Trp Leu Ala 65 70 75 80

Glu Gln Arg Val Pro Pro Ser Val Gln Leu Lys Pro Glu Val Ser Pro 85 90 95

Thr Gln Asp Ile Arg Leu Trp Val Ser Val Glu Asp Ala Gln Met His
100 105 110

Thr Val Thr Ile Trp Leu Thr Val Arg Pro Asp Met Thr Val Ala Ser 115 120 125 Leu Lys Asp Met Val Phe Leu Asp Tyr Gly Phe Pro Pro Val Leu Gln Gln Trp Val Ile Gly Gln Arg Leu Ala Arg Asp Gln Glu Thr Leu His Ser His Gly Val Arg Gln Asn Gly Asp Ser Ala Tyr Leu Tyr Leu Leu Ser Ala Arg Asn Thr Ser Leu Asn Pro Gln Glu Leu Gln Arg Glu Arg Gln Leu Arg Met Leu Glu Asp Leu Gly Phe Lys Asp Leu Thr Leu Gln Pro Arg Gly Pro Leu Glu Pro Gly Pro Pro Lys Pro Gly Val Pro Gln Glu Pro Gly Arg Gly Gln Pro Asp Ala Val Pro Glu Pro Pro Pro Val Gly Trp Gln Cys Pro Gly Cys Thr Phe Ile Asn Lys Pro Thr Arg Pro Gly Cys Glu Met Cys Cys Arq Ala Arg Pro Glu Ala Tyr Gln Val Pro Ala Ser Tyr Gln Pro Asp Glu Glu Glu Arg Ala Arg Leu Ala Gly Glu Glu Glu Ala Leu Arg Gln Tyr Gln Gln Arg Lys Gln Gln Gln Gln Glu Gly Asn Tyr Leu Gln His Val Gln Leu Asp Gln Arg Ser Leu Val Leu Asn Thr Glu Pro Ala Glu Cys Pro Val Cys Tyr Ser Val Leu Ala Pro Gly Glu Ala Val Leu Arg Glu Cys Leu His Thr Phe Cys Arg Glu Cys Leu Gln Gly Thr Ile Arg Asn Ser Gln Glu Ala Glu Val Ser Cys Pro Phe Ile Asp Asn Thr Tyr Ser Cys Ser Gly Lys Leu Leu Glu Arg

Glu Ile Lys Ala Leu Leu Thr Pro Glu Asp Tyr Gln Arg Phe Leu Asp 385 390 Leu Gly Ile Ser Ile Ala Glu Asn Arg Ser Ala Phe Ser Tyr His Cys 405 410 415 Lys Thr Pro Asp Cys Lys Gly Trp Cys Phe Phe Glu Asp Asp Val Asn 420 Glu Phe Thr Cys Pro Val Cys Phe His Val Asn Cys Leu Leu Cys Lys 435 440 445 Ala Ile His Glu Gln Met Asn Cys Lys Glu Tyr Gln Glu Asp Leu Ala 450 455 460 Leu Arg Ala Gln Asn Asp Val Ala Ala Arg Gln Thr Thr Glu Met Leu 465 470 475 480 Lys Val Met Leu Gln Gln Gly Glu Ala Met Arg Cys Pro Gln Cys Gln 485 490 495 Ile Val Val Gln Lys Lys Asp Gly Cys Asp Trp Ile Arg Cys Thr Val 500 505 510 Cys His Thr Glu Ile Cys Trp Val Thr Lys Gly Pro Arg Trp Gly Pro 515 520 525 Gly Gly Pro Gly Asp Thr Ser Gly Gly Cys Arg Cys Arg Val Asn Gly 530 535 540 Ile Pro Cys His Pro Ser Cys Gln Asn Cys His 545 550 <210> 37 <211> 1233 <212> DNA <213> Homo sapiens <400> 37 agacgtggga tgcacacagc tcagaacagt tggatcttgc tcagtctctg tcagaggaag 60 atcccttgga caagaggacc ctgccttggt gtgagagtga gggaagagga agctggaacg 120 agggttaagg aaaaccttcc agtctggaca gtgactggag agctccaagg aaagcccctc 180 ggtaacccag ccgctggcac catgaaccca gagagcagta tctttattqa qqattacctt 240

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<212> PRT

<213> Homo sapiens

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Thr Gly Glu Leu Gln Gly Lys Pro Leu Gly Asn Pro Ala Ala Gly Thr 50 55 60

Met Asn Pro Glu Ser Ser Ile Phe Ile Glu Asp Tyr Leu Lys Tyr Phe 65 70 75 80

Gln Asp Gln Val Ser Arg Glu Asn Leu Leu Gln Leu Leu Thr Asp Asp 85 90 95

Glu Ala Trp Asn Gly Phe Val Ala Ala Ala Glu Leu Pro Arg Asp Glu 100 105 110

Ala Asp Glu Leu Arg Lys Ala Leu Asn Lys Leu Ala Ser His Met Val 115 120 125

Met Lys Asp Lys Asn Arg His Asp Lys Asp Gln Gln His Arg Gln Trp 130 135 140

| Phe 145 | Leu | Lys | Glu | Phe | Pro 150 | Arg | Leu | Lys | Arg | Glu 155 | Leu | Glu | Asp | His | Ile 160 |
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| Arg | Lys | Leu | Arg | Ala 165 | Leu | Ala | Glu | Glu | Val 170 | Glu | Gln | Val | His | Arg 175 | Gly |
| Thr | Thr | Ile | Ala 180 | Asn | Val | Val | Ser | Asn 185 | Ser | Val | Gly | Thr | Thr 190 | Ser | Gly |
| Ile | Leu | Thr 195 | Leu | Leu | Gly | Leu | Gly 200 | Leu | Ala | Pro | Phe | Thr 205 | Glu | Gly | Ile |
| Ser | Phe 210 | Val | Leu | Leu | Asp | Thr 215 | Gly | Met | Gly | Leu | Gly 220 | Ala | Ala | Ala | Ala |
| Val 225 | Ala | Gly | Ile | Thr | Cys 230 | Ser | Val | Val | Glu | Leu 235 | Val | Asn | Lys | Leu | Arg 240 |
| Ala | Arg | Ala | Gln | Ala 245 | Arg | Asn | Leu | Asp | Gln 250 | Ser | Gly | Thr | Asn | Val 255 | Ala |
| Lys | Val | Met | Lys 260 | Glu | Phe | Val | Gly | Gly 265 | Asn | Thr | Pro | Asn | Val 270 | Leu | Thr |
| Leu | Val | Asp 275 | Asn | Trp | Tyr | Gln | Val 280 | Thr | Gln | Gly | Ile | Gly 285 | Arg | Asn | Ile |
| Arg | Ala 290 | Ile | Arg | Arg | Ala | Arg 295 | Ala | Asn | Pro | Gln | Leu 300 | Gly | Ala | Tyr | Ala |
| Pro 305 | Pro | Pro | His | Val | Ile 310 | Gly | Arg | Ile | Ser | Ala 315 | Glu | Gly | Gly | Glu | Gln 320 |
| Val | Glu | Arg | Val | Val 325 | Glu | Gly | Pro | Ala | Gln 330 | Ala | Met | Ser | Arg | Gly 335 | Thr |
| Met | Ile | Val | Gly 340 | Ala | Ala | Thr | Gly | Gly 345 | Ile | Leu | Leu | Leu | Leu 350 | Asp | Val |
| Val | Ser | Leu 355 | Ala | Tyr | Glu | Ser | Lys 360 | His | Leu | Leu | Glu | Gly 365 | Ala | Lys | Ser |
| Glu | Ser 370 | Ala | Glu | Glu | Leu | Lys 375 | Lys | Arg | Ala | Gln | Glu 380 | Leu | Glu | Gly | Lys |
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cctggaatgg attcgtggct gctgctgaac tgcccaggga tgaggcagat gagctccgta 360
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accagcagca caggcagtgg tttttgaaag agtttcctcg gttgaaaagg gagcttgagg 480
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Lys Ile Pro Trp Thr Arg Gly Pro Cys Leu Gly Val Arg Val Arg Glu
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                                 25
                                                      30
Glu Glu Ala Gly Thr Arg Val Lys Glu Asn Leu Pro Val Trp Thr Val
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                             40
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| Ί | hr | Gly 50 | Glu | Leu | Gln | Gly | Lys 55 | Pro | Leu | Gly | Asn | Pro 60 | Ala | Ala | Gly | Thr |
|---|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| M | let 65 | Asn | Pro | Glu | Ser | Ser 70 | Ile | Phe | Ile | Glu | Asp 75 | _ | Leu | Lys | Tyr | Phe 80 |
| (| Sln | Asp | Gln | Val | Ser 85 | Arg | Glu | Asn | Leu | Leu 90 | Gln | Leu | Leu | Thr | Asp 95 | Asp |
| (| Slu | Ala | Trp | Asn 100 | Gly | Phe | Val | Ala | Ala 105 | Ala | Glu | Leu | Pro | Arg 110 | Asp | Glu |
| P | Ala | Asp | Glu 115 | Leu | Arg | Lys | Ala | Leu 120 | Asn | Lys | Leu | Ala | Ser 125 | His | Met | Val |
| M | let | Lys 130 | Asp | Lys | Asn | Arg | His 135 | Asp | Lys | Asp | Gln | Gln 140 | His | Arg | Gln | Trp |
| | he .45 | Leu | Lys | Glu | Phe | Pro 150 | Arg | Leu | Lys | Arg | Glu 155 | Leu | Glu | Asp | His | Ile 160 |
| P | Arg | Lys | Leu | Arg | Ala 165 | Leu | Ala | Glu | Glu | Val 170 | Glu | Gln | Val | His | Arg 175 | Gly |
| Γ | hr. | Thr | Ile | Ala 180 | Asn | Val | Val | Ser | Asn 185 | Ser | Val | Gly | Thr | Thr 190 | Ser | Gly |
| Ι | le | Leu | Thr 195 | Leu | Leu | Gly | Leu | Gly 200 | Leu | Ala | Pro | Phe | Thr 205 | Glu | Gly | Ile |
| S | Ser | Phe 210 | Val | Leu | Leu | Asp | Thr 215 | Gly | Met | Gly | Leu | Gly 220 | Ala | Ala | Ala | Ala |
| | 7al 225 | Ala | Gly | Ile | Thr | Cys 230 | Ser | Val | Val | Glu | Leu 235 | Val | Asn | Lys | Leu | Arg 240 |
| P | Ala | Arg | Ala | Gln | Ala 245 | Arg | Asn | Leu | Asp | Gln 250 | Ser | Gly | Thr | Asn | Val 255 | Ala |
| I | ys | Val | Met | Lys 260 | Glu | Phe | Val | Gly | Gly 265 | Asn | Thr | Pro | Asn | Val 270 | Leu | Thr |
| I | eu | Val | Asp 275 | Asn | Trp | Tyr | Gln | Val 280 | Thr | Gln | Gly | Ile | Gly 285 | Arg | Asn | Ile |
| P | rg | Ala | Ile | Arg | Arg | Ala | Arg | Ala | Asn | Pro | Gln | Leu | Gly | Ala | Tyr | Ala |

Pro Pro Pro His Val Ile Gly Arg Ile Ser Ala Glu Gly Gly Glu Gln 305 310 315 320

Val Glu Arg Val Val Glu Gly Pro Ala Gln Ala Met Ser Arg Gly Thr 325 330 335

Met Ile Val Gly Ala Ala Thr Gly Gly Ile Leu Leu Leu Leu Asp Val 340 345 350

Val Ser Leu Ala Tyr Glu Ser Lys His Leu Leu Glu Gly Ala Lys Ser 355 360 365

Glu Ser Ala Glu Glu Leu Lys Lys Arg Ala Gln Glu Leu Glu Gly Lys 370 375 380

Leu Asn Phe Leu Thr Lys Ile His Glu Met Leu Gln Pro Gly Gln Asp 385 390 395 400

Gln

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Asp Phe Glu Ala Asp Ala His Trp Trp Ser Glu Arg Thr His Lys Asn
35 40 45

Leu Ser Asp Met Glu Asn Glu Phe Tyr Tyr Arg Tyr Pro Ser Phe Gln 50 55 60

Asp Val His Val Met Val Phe Val Gly Phe Gly Phe Leu Met Thr Phe 65 70 75 80

Leu Gln Arg Tyr Gly Phe Ser Ala Val Gly Phe Asn Phe Leu Leu Ala 85 90 95

Ala Phe Gly Ile Gln Trp Ala Leu Leu Met Gln Gly Trp Phe His Phe 100 105 110

Leu Gln Asp Arg Tyr Ile Val Val Gly Val Glu Asn Leu Ile Asn Ala 115 120 125

Asp Phe Cys Val Ala Ser Val Cys Val Ala Phe Gly Ala Val Leu Gly 130 135 140

Lys Val Ser Pro Ile Gln Leu Leu Ile Met Thr Phe Phe Gln Val Thr 145 150 155 160

Leu Phe Ala Val Asn Glu Phe Ile Leu Leu Asn Leu Leu Lys Val Lys 165 170 175

Asp Ala Gly Gly Ser Met Thr Ile His Thr Phe Gly Ala Tyr Phe Gly 180 185 190

Leu Thr Val Thr Arg Ile Leu Tyr Arg Arg Asn Leu Glu Gln Ser Lys

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| Thr 225 | Leu | Phe | Leu | Trp | Met 230 | Tyr | Trp | Pro | Ser | Phe 235 | Asn | Ser | Ala | Ile | Ser 240 |
| Tyr | His | Gly | Asp | Ser 245 | Gln | His | Arg | Ala | Ala 250 | Ile | Asn | Thr | Tyr | Cys 255 | Ser |
| Leu | Ala | Ala | Cys 260 | Val | Leu | Thr | Ser | Val 265 | Ala | Ile | Ser | Ser | Ala 270 | Leu | His |
| Lys | Lys | Gly 275 | Lys | Leu | Asp | Met | Val 280 | His | Ile | Gln | Asn | Ala 285 | Thr | Leu | Ala |
| Gly | Gly 290 | Val | Ala | Val | Gly | Thr 295 | Ala | Ala | Glu | Met | Met 300 | Leu | Met | Pro | Tyr |
| Gly 305 | Ala | Leu | Ile | Ile | Gly 310 | Phe | Val | Cys | Gly | Ile 315 | Ile | Ser | Thr | Leu | Gly 320 |
| Phe | Val | Tyr | Leu | Thr 325 | Pro | Phe | Leu | Glu | Ser 330 | Arg | Leu | His | Ile | Gln 335 | Asp |
| Thr | Cys | Gly | Ile 340 | Asn | Asn | Leu | His | Gly 345 | Ile | Pro | Gly | Ile | Ile 350 | Gly | Gly |
| Ile | Val | Gly 355 | Ala | Val | Thr | Ala | Ala 360 | Ser | Ala | Ser | Leu | Glu 365 | Val | Tyr | Gly |
| Lys | Glu 370 | Gly | Leu | Val | His | Ser 375 | Phe | Asp | Phe | Gln | Gly 380 | Phe | Asn | Gly | Asp |
| Trp 385 | Thr | Ala | Arg | Thr | Gln 390 | Gly | Lys | Phe | Gln | Ile 395 | Tyr | Gly | Leu | Leu | Val 400 |
| Thr | Leu | Ala | Met | Ala 405 | Leu | Met | Gly | Gly | Ile 410 | Ile | Val | Gly | Leu | Ile 415 | Leu |
| Arg | Leu | Pro | Phe 420 | Trp | Gly | Gln | Pro | Ser 425 | Asp | Glu | Asn | Cys | Phe 430 | Glu | Asp |
| Ala | Val | Tyr 435 | Trp | Glu | Val | Ser | Ser 440 | Arg | Asp | Leu | Ala | Pro 445 | | | |

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<400> 44
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| | | 2 5 | | | | | 4.0 | | | | | 4 5 | | | |

Cys

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- Glu Thr Ala Ile Asn Asn Leu Asn Pro Ala Phe Ser Lys Lys Phe Val
- Leu Asp Tyr His Phe Glu Glu Val Gln Lys Leu Lys Phe Ala Leu Phe
- Asp Gln Asp Lys Ser Ser Met Arg Leu Asp Glu His Asp Phe Leu Gly
- Gln Phe Ser Cys Ser Leu Gly Thr Ile Val Ser Ser Lys Lys Ile Thr
- Arg Pro Leu Leu Leu Asn Asp Lys Pro Ala Gly Lys Gly Leu Ile
- Thr Ile Ala Ala Gln Glu Leu Ser Asp Asn Arg Val Ile Thr Leu Ser
- Leu Ala Gly Arg Arg Leu Asp Lys Lys Asp Leu Phe Gly Lys Ser Asp
- Pro Phe Leu Glu Phe Tyr Lys Pro Gly Asp Asp Gly Lys Trp Met Leu
- Val His Arg Thr Glu Val Ile Lys Tyr Thr Leu Asp Pro Val Trp Lys
- Pro Phe Thr Val Pro Leu Val Ser Leu Cys Asp Gly Asp Met Glu Lys
- Pro Ile Gln Val Met Cys Tyr Asp Tyr Asp Asn Asp Gly His Asp
- Phe Ile Gly Glu Phe Gln Thr Ser Val Ser Gln Met Cys Glu Ala Arg
- Asp Ser Val Pro Leu Glu Phe Glu Cys Ile Asn Pro Lys Lys Gln Arg
- Lys Lys Lys Asn Tyr Lys Asn Ser Gly Ile Ile Leu Arg Ser Cys

Lys Ile Asn Arg Asp Tyr Ser Phe Leu Asp Tyr Ile Leu Gly Gly Cys Gln Leu Met Phe Thr Val Gly Ile Asp Phe Thr Ala Ser Asn Gly Asn Pro Leu Asp Pro Ser Ser Leu His Tyr Ile Asn Pro Met Gly Thr Asn Glu Tyr Leu Ser Ala Ile Trp Ala Val Gly Gln Ile Ile Gln Asp Tyr Asp Ser Asp Lys Met Phe Pro Ala Leu Gly Phe Gly Ala Gln Leu Pro Pro Asp Trp Lys Val Ser His Glu Phe Ala Ile Asn Phe Asn Pro Thr Asn Pro Phe Cys Ser Gly Val Asp Gly Ile Ala Gln Ala Tyr Ser Ala Cys Leu Pro His Ile Arg Phe Tyr Gly Pro Thr Asn Phe Ser Pro Ile Val Asn His Val Ala Arg Phe Ala Ala Gln Ala Thr Gln Gln Arg Thr Ala Thr Gln Tyr Phe Ile Leu Leu Ile Ile Thr Asp Gly Val Ile Ser Asp Met Glu Glu Thr Arg His Ala Val Val Gln Ala Ser Lys Leu Pro Met Ser Ile Ile Ile Val Gly Val Gly Asn Ala Asp Phe Ala Ala Met Glu Phe Leu Asp Gly Asp Ser Arg Met Leu Arg Ser His Thr Gly Glu Glu Ala Ala Arg Asp Ile Val Gln Phe Val Pro Phe Arg Glu Phe Arg

Asn Ala Ala Lys Glu Thr Leu Ala Lys Ala Val Leu Ala Glu Leu Pro

Gln Gln Val Val Gln Tyr Phe Lys His Lys Asn Leu Pro Pro Thr Ser

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Tyr Glu Asn Pro Thr 545

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<211> 1070

<212> DNA

<213> Homo sapiens

<400> 45

atgttggcac tettggttet ggtgaetgtg geeetggeat etgeteatea tggtggtgag 60 cactttgaag gggagaaggt gttccgtgtt aacgttgaag atgaaaatca cattaacata 120 atcogcgagt tggccacctt tattcagatt gacttctgga agccagattc tgtcacacaa 180 atcaaacctc acagtacagt tgacttccgt gttaaagcag aagatactgt cactgtggag 240 aatgttctaa agcagaatga actacaatac aaggtactga taagcaacct gagaaatgtg 300 gtggaggctc agtttgatag ccgggttcgt qcaacaggac acagttatga gaagtacaac 360 aagtgggaaa cgatagaggc ttggactcaa caagtcgcca ctgagaatcc agccctcatc 420 tctcgcagtg ttatcggaac cacatttgag ggacgcgcta tttacctcct gaaggttggc 480 aaagctggac aaaataagcc tgccattttc atggaatgtg gtttccatgc cagagagtgg 540 atttctcctg cattctqcca qtqqtttqta aqaqaqqctq ttcqtaccta tqqacqtqaq 600 atccaagtga cagagettet egacaagtta gaettttatg teetgeetgt geteaatatt 660 gatggctaca tctacacctg gaccaagagc cgattttgga gaaagacttc gctccaccca 720 tactggatct accettactc atatgettac aaacteggtg agaacaatge tgagttgaat 780 gccctggcta aagctactgt gaaagaactt gcctcactgc acggcaccaa gtacacatat 840 ggcccgggag ctacaacaat ctatcctgct gctgggggct ctgacgactg ggcttatgac 900 caaggaatca gatatteett caeetttgaa ettegagata caggeagata tggetttete 960 cttccagaat cccagatccg ggctacctgc gaggagacct tcctggcaat caagtatgtt 1020 gccagctacg tcctggaaca cctgtactag ttgagaaagc tgatggcctt 1070

<210> 46

<211> 349

<212> PRT

<213> Homo sapiens

<400> 46

Met Leu Ala Leu Leu Val Leu Val Thr Val Ala Leu Ala Ser Ala His

1 5 10 15

His Gly Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg Val Asn Val 20 25 30

Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala Thr Phe Ile 35 40 45

Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile Lys Pro His

| 50 | 55 | 60 |
|----|----|----|
| | | |

| Ser 65 | Thr | Val | Asp | Phe | Arg 70 | Val | Lys | Ala | Glu | Asp 75 | Thr | Val | Thr | Val | Glu 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Asn | Val | Leu | Lys | Gln 85 | Asn | Glu | Leu | Gln | Tyr 90 | Lys | Val | Leu | Ile | Ser 95 | Asn |
| Leu | Arg | Asn | Val 100 | Val | Glu | Ala | Gln | Phe 105 | Asp | Ser | Arg | Val | Arg 110 | Ala | Thr |
| Gly | His | Ser 115 | Tyr | Glu | Lys | Tyr | Asn 120 | Lys | Trp | Glu | Thr | Ile 125 | Glu | Ala | Trp |
| Thr | Gln 130 | Gln | Val | Ala | Thr | Glu 135 | Asn | Pro | Ala | Leu | Ile 140 | Ser | Arg | Ser | Val |
| Ile 145 | Gly | Thr | Thr | Phe | Glu 150 | Gly | Arg | Ala | Ile | Tyr 155 | Leu | Leu | Lys | Val | Gly 160 |
| Lys | Ala | Gly | Gln | Asn 165 | Lys | Pro | Ala | Ile | Phe 170 | Met | Glu | Cys | Gly | Phe 175 | His |
| Ala | Arg | Glu | Trp 180 | Ile | Ser | Pro | Ala | Phe 185 | Cys | Gln | Trp | Phe | Val 190 | Arg | Glu |
| Ala | Val | Arg 195 | Thr | Tyr | Gly | Arg | Glu 200 | Ile | Gln | Val | Thr | Glu 205 | Leu | Leu | Asp |
| Lys | Leu 210 | Asp | Phe | Tyr | Val | Leu 215 | Pro | Val | Leu | Asn | Ile 220 | Asp | Gly | Tyr | Ile |
| Tyr 225 | Thr | Trp | Thr | Lys | Ser 230 | Arg | Phe | Trp | Arg | Lys 235 | Thr | Ser | Leu | His | Pro |
| Tyr | Trp | Ile | Tyr | Pro 245 | Tyr | Ser | Tyr | Ala | Tyr 250 | Lys | Leu | Gly | Glu | Asn 255 | Asn |
| Ala | Glu | Leu | Asn 260 | Ala | Leu | Ala | Lys | Ala 265 | Thr | Val | Lys | Glu | Leu 270 | Ala | Ser |
| Leu | His | Gly 275 | Thr | Lys | Tyr | Thr | Tyr 280 | Gly | Pro | Gly | Ala | Thr 285 | Thr | Ile | Tyr |
| Pro | Ala 290 | Ala | Gly | Gly | Ser | Asp 295 | Asp | Trp | Ala | Tyr | Asp 300 | Gln | Gly | Ile | Arg |

Tyr Ser Phe Thr Phe Glu Leu Arg Asp Thr Gly Arg Tyr Gly Phe Leu

Leu Pro Glu Ser Gln Ile Arg Ala Thr Cys Glu Glu Thr Phe Leu Ala 325 330 335

Ile Lys Tyr Val Ala Ser Tyr Val Leu Glu His Leu Tyr 340 345

<210> 47

<211> 693

<212> DNA

<213> Homo sapiens

<400> 47

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<210> 48

<211> 231

<212> PRT

<213> Homo sapiens

<400> 48

Gly Ser His His Gly Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg
1 5 10 15

Val Asn Val Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala 20 25 30

Ser Thr Thr Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile 35 40 45

Lys Pro His Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val 50 55 60

Thr Val Glu Asn Val Leu Lys Gln Asn Glu Leu Gln Tyr Lys Val Leu

65 70 75 80

Ile Ser Asn Leu Arg Asn Val Val Glu Ala Gln Phe Asp Ser Arg Val 85 90 95

Arg Ala Thr Gly His Ser Tyr Glu Lys Tyr Asn Lys Trp Glu Thr Ile 100 105 110

Glu Ala Trp Thr Gln Gln Val Ala Thr Glu Asn Pro Ala Leu Ile Ser 115 120 125

Arg Ser Val Ile Gly Thr Thr Phe Glu Gly Arg Ala Ile Tyr Leu Leu 130 135 140

Lys Val Gly Lys Ala Gly Gln Asn Lys Pro Ala Ile Phe Met Glu Cys 145 150 155 160

Gly Phe His Ala Arg Glu Trp Ile Ser Pro Ala Phe Cys Gln Trp Phe 165 170 175

Val Arg Glu Ala Val Arg Thr Tyr Gly Arg Glu Ile Gln Val Thr Glu 180 185 190

Leu Leu Asp Lys Leu Asp Phe Tyr Val Leu Pro Val Leu Asn Ile Asp 195 200 205

Gly Tyr Ile Tyr Thr Trp Thr Lys Ser Arg Phe Trp Arg Lys Thr Ser 210 215 220

Leu His Pro Tyr Trp Leu Glu 225 230

<210> 49

<211> 693

<212> DNA

<213> Homo sapiens

<400> 49

ggatcccatc atggtggtga gcactttgaa ggcgagaagg tgttccgtgt taacgttgaa 60 gatgaaaatc acattaacat aatccgcgag ttggccagca cgacccagat tgacttctgg 120 aagccagatt ctgtcacaca aatcaaacct cacagtacag ttgacttccg tgttaaagca 180 gaagatactg tcactgtgga gaatgttcta aagcagaatg aactacaata caaggtactg 240 ataagcaacc tgagaaatgt ggtggaggct cagtttgata gccgggttcg tgcaacagga 300 cacagttatg agaagtacaa caagtgggaa acgatagagg cttggactca acaagtcgcc 360 actgagaatc cagcctcat ctctcgcagt gttatcggaa ccacatttga gggacgcgct 420 atttacctcc tgaaggttgg caaagctgga caaaataagc ctgccatttt catggactgt 480 ggtttccatg ccagagagtg gatttctcct gcattctgcc agtggtttgt aagagaggct 540

gttcgtacca atggacgtga gatccaagtg acagagette tegacaagtt agaetttat 600 gteetgeetg tgeteaatat tgatggetae atetacaeet ggaccaagag eegattttgg 660 agaaagaett egeteeaee atactggete gag 693

<210> 50

<211> 231

<212> PRT

<213> Homo sapiens

<400> 50

Gly Ser His His Gly Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg
1 5 10 15

Val Asn Val Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala 20 25 30

Ser Thr Thr Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile 35 40 45

Lys Pro His Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val 50 55 60

Thr Val Glu Asn Val Leu Lys Gln Asn Glu Leu Gln Tyr Lys Val Leu 65 70 75 80

Ile Ser Asn Leu Arg Asn Val Val Glu Ala Gln Phe Asp Ser Arg Val
85 90 95

Arg Ala Thr Gly His Ser Tyr Glu Lys Tyr Asn Lys Trp Glu Thr Ile 100 105 110

Glu Ala Trp Thr Gln Gln Val Ala Thr Glu Asn Pro Ala Leu Ile Ser 115 120 125

Arg Ser Val Ile Gly Thr Thr Phe Glu Gly Arg Ala Ile Tyr Leu Leu 130 135 140

Lys Val Gly Lys Ala Gly Gln Asn Lys Pro Ala Ile Phe Met Asp Cys 145 150 155 160

Gly Phe His Ala Arg Glu Trp Ile Ser Pro Ala Phe Cys Gln Trp Phe 165 170 175

Val Arg Glu Ala Val Arg Thr Asn Gly Arg Glu Ile Gln Val Thr Glu 180 185 190

Leu Leu Asp Lys Leu Asp Phe Tyr Val Leu Pro Val Leu Asn Ile Asp

195 200 205

Gly Tyr Ile Tyr Thr Trp Thr Lys Ser Arg Phe Trp Arg Lys Thr Ser 210 215 220

Leu His Pro Tyr Trp Leu Glu 225 230

<210> 51

<211> 693

<212> DNA

<213> Homo sapiens

<400> 51

ggatcccatc atggtggtga gcactttgaa ggcgagaagg tgttccgtgt taacgttgaa 60 gatgaaaatc acattaacat aatccgcgag ttggccagca cgacccagat tgacttctgg 120 aagccagatt ctgtcacaca aatcaaacct cacagtacag ttgacttccg tgttaaagca 180 gaagatactg tcactgtgga gaatgttcta aagcagaatg aactacaata caaggtactg 240 ataagcaacc tgagaaatgt ggtggaggct cagtttgata gccgggttcg tgcaacagga 300 cacagttatg agaagtacaa caagtgggaa acgatagagg cttggactca acaagtcgcc 360 actgagaatc cagcctcat ctctcgcagt gttatcggaa ccacatttga gggacgcgct 420 atttacctcc tgaaggttgg caaagctgga caaaataagc ctgccattt catggactgt 480 ggttccatg ccagagagtg gatttctcct gcattctgcc agtggttgt aagaagaggct 540 gtccgcctg tgctcaatat tgatggctac atctacacct ggaccaagag ccgattttg 660 agaaagactt cgctcaccc atactggctc gag

<210> 52

<211> 231

<212> PRT

<213> Homo sapiens

<400> 52

Gly Ser His His Gly Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Val Asn Val Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala 20 25 30

Ser Thr Thr Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile 35 40 45

Lys Pro His Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val 50 55 60

Thr Val Glu Asn Val Leu Lys Gln Asn Glu Leu Gln Tyr Lys Val Leu

Ile Ser Asn Leu Arg Asn Val Val Glu Ala Gln Phe Asp Ser Arg Val 85 90 95

Arg Ala Thr Gly His Ser Tyr Glu Lys Tyr Asn Lys Trp Glu Thr Ile 100 105 110

Glu Ala Trp Thr Gln Gln Val Ala Thr Glu Asn Pro Ala Leu Ile Ser 115 120 125

Arg Ser Val Ile Gly Thr Thr Phe Glu Gly Arg Ala Ile Tyr Leu Leu 130 135 140

Lys Val Gly Lys Ala Gly Gln Asn Lys Pro Ala Ile Phe Met Asp Cys 145 150 155 160

Gly Phe His Ala Arg Glu Trp Ile Ser Pro Ala Phe Cys Gln Trp Phe 165 170 175

Val Arg Glu Ala Val Arg Thr Tyr Gly Arg Glu Ile Gln Val Thr Glu 180 185 190

Leu Leu Asp Lys Leu Asp Phe Tyr Val Leu Pro Val Leu Asn Ile Asp 195 200 205

Gly Tyr Ile Tyr Thr Trp Thr Lys Ser Arg Phe Trp Arg Lys Thr Ser 210 215 220

Leu His Pro Tyr Trp Leu Glu 225 230

<210> 53

<211> 693

<212> DNA

<213> Homo sapiens

<400> 53

ggatcccatc atggtggtga gcactttgaa ggcgagaagg tgttccgtgt taacgttgaa 60 gatgaaaatc acattaacat aatccgcgag ttggccagca cgacccagat tgacttctgg 120 aagccagatt ctgtcacaca aatcaaacct cacagtacag ttgacttccg tgttaaagca 180 gaagatactg tcactgtgga gaatgttcta aagcagaatg aactacaata caaggtactg 240 ataagcaacc tgagaaatgt ggtggaggct cagtttgata gccgggttcg tgcaacagga 300 cacagttatg agaagtacaa caagtgggaa acgatagagg cttggactca acaagtcgcc 360 actgagaatc cagcctcat ctctcgcagt gttatcggaa ccacatttga gggacgcgtt 420 atttacctcc tgaaggttgg caaagctgga caaaataagc ctgccatttt catggactgt 480 ggtttccatg ccagaagtg gatttctcct gcattccgcc agtggtttgt aagagaggct 540

gttcgtacct atggacgtga gatccaagtg acagagcttc tcgacaagtt agacttttat 600 gtcctgcctg tgctcaatat tgatggctac atctacacct ggaccaagag ccgattttgg 660 agaaagactt cgctccaccc atactggctc gag 693

<210> 54

<211> 231

<212> PRT

<213> Homo sapiens

<400> 54

Gly Ser His His Gly Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg

1 5 10 15

Val Asn Val Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala 20 25 30

Ser Thr Thr Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile 35 40 45

Lys Pro His Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val 50 55 60

Thr Val Glu Asn Val Leu Lys Gln Asn Glu Leu Gln Tyr Lys Val Leu 65 70 75 80

Ile Ser Asn Leu Arg Asn Val Val Glu Ala Gln Phe Asp Ser Arg Val 85 90 95

Arg Ala Thr Gly His Ser Tyr Glu Lys Tyr Asn Lys Trp Glu Thr Ile 100 105 110

Glu Ala Trp Thr Gln Gln Val Ala Thr Glu Asn Pro Ala Leu Ile Ser 115 120 125

Arg Ser Val Ile Gly Thr Thr Phe Glu Gly Arg Val Ile Tyr Leu Leu 130 135 140

Lys Val Gly Lys Ala Gly Gln Asn Lys Pro Ala Ile Phe Met Asp Cys 145 150 155 160

Gly Phe His Ala Arg Glu Trp Ile Ser Pro Ala Phe Arg Gln Trp Phe 165 170 175

Val Arg Glu Ala Val Arg Thr Tyr Gly Arg Glu Ile Gln Val Thr Glu 180 185 190

Leu Leu Asp Lys Leu Asp Phe Tyr Val Leu Pro Val Leu Asn Ile Asp

195 200 205

Gly Tyr Ile Tyr Thr Trp Thr Lys Ser Arg Phe Trp Arg Lys Thr Ser 210 215 220

Leu His Pro Tyr Trp Leu Glu 225 230

<210> 55

<211> 649

<212> DNA

<213> Homo sapiens

<400> 55

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<210> 56

<211> 161

<212> PRT

<213> Homo sapiens

<400> 56

Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp

1 5 10 15

His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu 20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Arg Phe Ala Lys Lys 35 40 45

His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala 50 55 60

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asp Arg Leu 85 90 95

Ala Tyr Ile Ala His Pro Lys Leu Gly Lys Arg Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys 115 120 125

Ala Lys Ala Lys Asp Gln Thr Lys Ala Gln Ala Ala Ala Pro Ala Ser 130 135 140

Val Pro Ala Gln Ala Pro Lys Arg Thr Gln Ala Pro Thr Lys Ala Ser 145 150 155 160

Glu

<210> 57

<211> 580

<212> DNA

<213> Homo sapiens

<400> 57

actcactata gggctcgagc ggcgcttcgg gagccgcggc ttatggtgca gacatggcca 60 agtccaagaa ccacaccaca cacaaccagt cccgaaaatg gcacagaaat ggtatcaaga 120 aaccccgatc acaaagatac gaatctctta agggggtgga ccccaagttc ctgaggaaca 180 tgcgctttgc caagaagcac aacaaaaagg gcctaaagaa gatgcaggcc aacaatgcca 240 aggccatgag tgcacgtgcc gaggctatca aggccctcgt aaagcccaag gaggttaagc 300 ccaagatccc aaagggtgtc agccgcaagc tcgatcgact tgcctacatt gcccacccca 360 agcttgggaa gcgtgctcgt gcccgtattg ccaaggggct caggctgtgc cggccaaagg 420 ccaaggccaa ggccaaagcc aaggccaagg atcaaaccaa ggcccaggct gcagcccag 480 cttcagttcc agctcaggc ccaaaaggc cccaaaggcc tacaaaggcc tcaaaaggcc tcaaaaggcc 580 tatctctgcc aacatgagga cagaaagact ggtgcgaccc

<210> 58

<211> 161

<212> PRT

<213> Homo sapiens

<400> 58

Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu

20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Arg Phe Ala Lys Lys 35 40 45

His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala 50 55 60

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asp Arg Leu 85 90 95

Ala Tyr Ile Ala His Pro Lys Leu Gly Lys Arg Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys 115 120 125

Ala Lys Ala Lys Asp Gln Thr Lys Ala Gln Ala Ala Pro Ala Ser 130 135 140

Val Pro Ala Gln Ala Pro Lys Arg Thr Gln Ala Pro Thr Lys Ala Ser 145 150 155 160

Glu

<210> 59

<211> 1143

<212> DNA

<213> Homo sapiens

<400> 59

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cactgcatgt atgggtggc acctccattc tgtgaggaag tggggtatgg aggaagcatt 780 gacagtgggc ctccaggact gctcagaggg gcgattcct cgtcaatttg ggttgtgtcc 840 atcataatgt ttcgccttat tttattaatc ctttcagtgg tttttgtgtt tttccggcaa 900 gtgataggaa accacttaaa acccaaacag gaaaaaatgc cactatccaa agcaaaaact 960 gaacaggaag aatctaaaac aaaaactgta caggaagaat ctaaaacaaa aactggacag 1020 gaagaatctg aagcaaaaac tggacaggaa gaatctaaag caaaaactgg acaggaagaa 1080 tctaaagcaa acattgaaag taaacgacc aaagcaaaga gtgtcaagaa acaaaaaaag 1140 taa

<210> 60

<211> 380

<212> PRT

<213> Homo sapiens

<400> 60

Met Arg Ser Val Gln Ile Phe Leu Ser Gln Cys Arg Leu Leu Leu 1 5 10 15

Leu Val Pro Thr Met Leu Leu Lys Ser Leu Gly Glu Asp Val Ile Phe 20 25 30

His Pro Glu Gly Glu Phe Asp Ser Tyr Glu Val Thr Ile Pro Glu Lys
35 40 45

Leu Ser Phe Arg Gly Glu Val Gln Gly Val Val Ser Pro Val Ser Tyr 50 55 60

Leu Leu Gln Leu Lys Gly Lys Lys His Val Leu His Leu Trp Pro Lys 65 70 75 80

Arg Leu Leu Pro Arg His Leu Arg Val Phe Ser Phe Thr Glu His
85 90 95

Gly Glu Leu Leu Glu Asp His Pro Tyr Ile Pro Lys Asp Cys Asn Tyr
100 105 110

Met Gly Ser Val Lys Glu Ser Leu Asp Ser Lys Ala Thr Ile Ser Thr 115 120 125

Cys Met Gly Gly Leu Arg Gly Val Phe Asn Ile Asp Ala Lys His Tyr 130 135 140

Leu Leu Lys Lys Glu Gln Phe Gly Asn Gln Ala Glu Asn Leu Met Cys 165 170 175

Trp Gly Thr Gly Tyr His Leu Ser Met Lys Pro Met Gly Ile Pro Asp Leu Gly Met Ile Asn Asp Gly Thr Ser Cys Gly Glu Gly Arg Val Cys Phe Lys Lys Asn Cys Val Asn Ser Ser Val Leu Gln Phe Asp Cys Leu Pro Glu Lys Cys Asn Thr Arg Gly Val Cys Asn Asn Arg Lys Ser Cys His Cys Met Tyr Gly Trp Ala Pro Pro Phe Cys Glu Glu Val Gly Tyr Gly Gly Ser Ile Asp Ser Gly Pro Pro Gly Leu Leu Arg Gly Ala Ile Pro Ser Ser Ile Trp Val Val Ser Ile Ile Met Phe Arq Leu Ile Leu Leu Ile Leu Ser Val Val Phe Val Phe Phe Arg Gln Val Ile Gly Asn His Leu Lys Pro Lys Gln Glu Lys Met Pro Leu Ser Lys Ala Lys Thr Glu Gln Glu Glu Ser Lys Thr Lys Thr Val Gln Glu Glu Ser Lys Thr Lys Thr Gly Gln Glu Glu Ser Glu Ala Lys Thr Gly Gln Glu Glu Ser Lys Ala Lys Thr Gly Gln Glu Glu Ser Lys Ala Asn Ile Glu Ser Lys Arg Pro Lys Ala Lys Ser Val Lys Lys Gln Lys Lys <210> 61

<211> 1207

<212> DNA

<213> Homo sapiens

<400> 61

ccgcgggact ccgccgtccc cqcccccag tcctccctcc cctccctcc agcatggtgc 60

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tcggggggga gcgctccagc cggccagccc cgtccgtggc gcccgagccg gacggctgcc 240
ccgtgtgcgt atggcggcag cacagccgcg agctgcgcct agagagcatc aagtcgcaga 300
tettgageaa actgeggete aaggaggege eeaacateag eegegaggtg gtgaageage 360
tgctgcccaa ggcgccgccg ctgcagcaga tcctggacct acacgacttc cagggcgacg 420
cgctgcagcc cgaggacttc ctggaggagg acgagtacca cqccaccacc qagaccqtca 480
ttagcatggc ccaggagacg gacccagcag tacagacaga tggcagccct ctctgctgcc 540
attiticacti cagococaag gigatgitica caaagagoat cgacticaag caagigotac 600
acagctggtt ccgccagcca cagagcaact ggggcatcga gatcaacgcc tttgatccca 660
gtggcacaga cctggctgtc acctccctgg ggccgggagc cgaggggctg catccattca 720
tggagcttcg agtcctagag aacacaaaac gttcccggcg gaacctgggt ctggactgcg 780
acgagcactc aagcgagtcc cgctgctgcc gatatcccct cacagtggac tttgaggctt 840
teggetggga etggateate geacetaage getacaagge caactaetge teeggeeagt 900
gcgagtacat gttcatgcaa aaatatccgc atacccattt ggtgcagcag gccaatccaa 960
gaggetetge tgggeeetgt tgtaceeeca ecaagatgte eccaateaac atgetetaet 1020
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gctgctctta agtgggtcac tacaagctgc tggagcaaag acttggtggg tgggtaactt 1140
aacctcttca cagaggataa aaaatgcttg tgagtatgac agaagggaat aaacaggctt 1200
aaagggt
                                                               1207
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<210> 62

<211> 345

<212> PRT

<213> Homo sapiens

<400> 62

Met Val Leu Ala Ala Pro Leu Leu Gly Phe Leu Leu Leu Ala Leu 1 5 10 15

Glu Leu Arg Pro Arg Gly Glu Ala Ala Glu Gly Pro Ala Ala Ala Ala 20 25 30

Ala Ala Ala Ala Ala Ala Ala Gly Val Gly Glu Arg Ser 35 40 45

Ser Arg Pro Ala Pro Ser Val Ala Pro Glu Pro Asp Gly Cys Pro Val 50 55 60

Cys Val Trp Arg Gln His Ser Arg Glu Leu Arg Leu Glu Ser Ile Lys
65 70 75 80

Ser Gln Ile Leu Ser Lys Leu Arg Leu Lys Glu Ala Pro Asn Ile Ser 85 90 95

Arg Glu Val Val Lys Gln Leu Leu Pro Lys Ala Pro Pro Leu Gln Gln
100 105 110

| Ile | Leu | Asp 115 | Leu | His | Asp | Phe | Gln 120 | Gly | Asp | Ala | Leu | Gln 125 | Pro | Glu | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Phe | Leu 130 | Glu | Glu | Asp | Glu | Tyr 135 | His | Ala | Thr | Thr | Glu 140 | Thr | Val | Ile | Ser |
| Met 145 | Ala | Gln | Glu | Thr | Asp 150 | Pro | Ala | Val | Gln | Thr 155 | Asp | Gly | Ser | Pro | Leu 160 |
| Cys | Cys | His | Phe | His 165 | Phe | Ser | Pro | Lys | Val 170 | Met | Phe | Thr | Lys | Ser 175 | Ile |
| Asp | Phe | Lys | Gln 180 | Val | Leu | His | Ser | Trp 185 | Phe | Arg | Gln | Pro | Gln 190 | Ser | Asn |
| Trp | Gly | Ile 195 | Glu | Ile | Asn | Ala | Phe 200 | Asp | Pro | Ser | Gly | Thr 205 | Asp | Leu | Ala |
| Val | Thr 210 | Ser | Leu | Gly | Pro | Gly 215 | Ala | Glu | Gly | Leu | His 220 | Pro | Phe | Met | Glu |
| Leu 225 | Arg | Val | Leu | Glu | Asn 230 | Thr | Lys | Arg | Ser | Arg 235 | Arg | Asn | Leu | Gly | Leu 240 |
| Asp | Cys | Asp | Glu | His 245 | Ser | Ser | Glu | Ser | Arg 250 | Cys | Cys | Arg | Tyr | Pro 255 | Leu |
| Thr | Val | Asp | Phe 260 | Glu | Ala | Phe | Gly | Trp 265 | Asp | Trp | Ile | Ile | Ala 270 | Pro | Lys |
| Arg | Tyr | Lys 275 | Ala | Asn | Tyr | Cys | Ser 280 | Gly | Gln | Cys | Glu | Tyr 285 | Met | Phe | Met |
| Gln | Lys 290 | Tyr | Pro | His | Thr | His 295 | Leu | Val | Gln | Gln | Ala 300 | Asn | Pro | Arg | Gly |
| Ser 305 | Ala | Gly | Pro | Cys | Cys 310 | Thr | Pro | Thr | Lys | Met 315 | Ser | Pro | Ile | Asn | Met 320 |
| Leu | Tyr | Phe | Asn | Asp 325 | Lys | Gln | Gln | Ile | Ile 330 | Tyr | Gly | Lys | Ile | Pro 335 | Gly |
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geetgetgea geageeeee accetgagee aageetgage ttteaggeae accatttget 1260
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<213> Homo sapiens
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Ala Val Pro Thr Ser Asp Leu Gly Glu Ile His Asn Trp Thr Glu Leu
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Leu Asp Leu Phe Asn His Thr Leu Ser Glu Cys His Val Glu Leu Ser
         35
                             40
                                                 45
Gln Ser Thr Lys Arg Val Val Leu Phe Ala Leu Tyr Leu Ala Met Phe
     50
                         55
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| Val 65 | Val | Gly | Leu | Val | Glu 70 | Asn | Leu | Leu | Val | Ile 75 | Cys | Val | Asn | Trp | Arg 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly | Ser | Gly | Arg | Ala 85 | Gly | Leu | Met | Asn | Leu 90 | Tyr | Ile | Leu | Asn | Met 95 | Ala |
| Ile | Ala | Asp | Leu 100 | Gly | Ile | Val | Leu | Ser 105 | Leu | Pro | Val | Trp | Met 110 | Leu | Glu |
| Val | Thr | Leu 115 | Asp | Tyr | Thr | Trp | Leu 120 | Trp | Gly | Ser | Phe | Ser 125 | Cys | Arg | Phe |
| Thr | His 130 | Tyr | Phe | Tyr | Phe | Val 135 | Asn | Met | Tyr | Ser | Ser 140 | Ile | Phe | Phe | Leu |
| Val 145 | Cys | Leu | Ser | Val | Asp 150 | Arg | Tyr | Val | Thr | Leu 155 | Thr | Ser | Ala | Ser | Pro 160 |
| Ser | Trp | Gln | Arg | Tyr 165 | Gln | His | Arg | Val | Arg 170 | Arg | Ala | Met | Cys | Ala 175 | Gly |
| Ile | Trp | Val | Leu 180 | Ser | Ala | Ile | Ile | Pro 185 | Leu | Pro | Glu | Val | Val 190 | His | Ile |
| Gln | Leu | Val 195 | Glu | Gly | Pro | Glu | Pro 200 | Met | Cys | Leu | Phe | Met 205 | Ala | Pro | Phe |
| Glu | Thr 210 | Tyr | Ser | Thr | Trp | Ala 215 | Leu | Ala | Val | Ala | Leu 220 | Ser | Thr | Thr | Ile |
| Leu 225 | Gly | Phe | Leu | Leu | Pro 230 | Phe | Pro | Leu | Ile | Thr 235 | Val | Phe | Asn | Val | Leu 240 |
| Thr | Ala | Cys | Arg | Leu 245 | Arg | Gln | Pro | Gly | Gln 250 | Pro | Lys | Ser | Arg | Arg 255 | His |
| Cys | Leu | Leu | Leu 260 | Cys | Ala | Tyr | Val | Ala 265 | Val | Phe | Val | Met | Cys 270 | Trp | Leu |
| Pro | Tyr | His 275 | Val | Thr | Leu | Leu | Leu 280 | Leu | Thr | Leu | His | Gly 285 | Thr | His | Ile |
| Ser | Leu 290 | His | Cys | His | Leu | Val 295 | His | Leu | Leu | Tyr | Phe 300 | Phe | Tyr | Asp | Val |
| Ile 305 | Asp | Cys | Phe | Ser | Met 310 | Leu | His | Суѕ | Val | Ile 315 | Asn | Pro | Ile | Leu | Tyr 320 |

Asn Phe Leu Ser Pro His Phe Arg Gly Arg Leu Leu Asn Ala Val Val 325 330 335

His Tyr Leu Pro Lys Asp Gln Thr Lys Ala Gly Thr Cys Ala Ser Ser 340 345 350

Ser Ser Cys Ser Thr Gln His Ser Ile Ile Ile Thr Lys Gly Asp Ser 355 360 365

Gln Pro Ala Ala Ala Pro His Pro Glu Pro Ser Leu Ser Phe Gln 370 375 380

Ala His His Leu Leu Pro Asn Thr Ser Pro Ile Ser Pro Thr Gln Pro 385 390 395 400

Leu Thr Pro Ser

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<211> 945

<212> DNA

<213> Homo sapiens

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<210> 66

<211> 314

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<213> Homo sapiens

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Val Lys Pro Ser Trp Gly Pro Gly Pro Ser Glu Gly Val Thr

Ala Val Pro Thr Ser Asp Leu Gly Glu Ile His Asn Trp Thr Glu Leu

Leu Asp His Leu Phe Asn His Thr Leu Ser Glu Cys His Val Glu Leu

Ser Gln Ser Thr Lys Arg Val Val Leu Phe Ala Leu Tyr Leu Ala Met

Phe Val Val Gly Leu Val Glu Asn Leu Leu Val Ile Cys Val Asn Trp

Arg Gly Ser Gly Arg Ala Gly Leu Met Asn Leu Tyr Ile Leu Asn Met

Ala Ile Ala Asp Leu Gly Ile Val Leu Ser Leu Pro Val Trp Met Pro

Glu Val Thr Leu Asp Tyr Thr Trp Leu Trp Gly Ser Phe Ser Cys Arg

Phe Thr His Tyr Phe Tyr Phe Val Asn Met Tyr Ser Ser Ile Phe Phe

Leu Val Cys Leu Ser Val Asp Arg Tyr Val Thr Leu Thr Gly Gln Pro

Lys Ser Arg Arg His Cys Leu Leu Cys Ala Tyr Val Ala Val Phe

Val Met Cys Trp Leu Pro Tyr His Val Thr Leu Leu Leu Leu Thr Leu

His Gly Thr His Ile Ser Leu His Cys His Leu Val His Leu Leu Tyr

Phe Phe Tyr Asp Val Ile Asp Cys Phe Ser Met Leu His Cys Val Ile

Asn Pro Ile Leu Tyr Asn Phe Leu Ser Pro His Phe Arg Gly Arg Leu

Leu Asn Ala Val Val His Tyr Leu Pro Lys Asp Gln Thr Lys Ala Gly

Thr Cys Ala Ser Ser Ser Cys Ser Thr Gln His Ser Ile Ile Ile 260 265 270 Thr Lys Gly Asp Ser Gln Pro Ala Ala Ala Ala Pro His Pro Glu 275 280 285 Pro Ser Leu Ser Phe Gln Ala His His Leu Leu Pro Asn Thr Ser Pro 290 295 300 Ile Ser Pro Thr Gln Pro Leu Thr Pro Ser 305 310 <210> 67 <211> 965 <212> DNA <213> Homo sapiens <400> 67 cgatgtcagt gaaacccagc tgggggcctg gccctcgga gggggtcacc gcagtgccta 60 ccagtgacct tggagagatc cacaactgga ccgagctgct tgacctcttc aaccacactt 120 tgtctgagtg ccacgtggag ctcagccaga gcaccaagcg cgtggtcctc tttgccctct 180 acctggccat gtttgtggtt gggctggtgg agaacctcct ggtgatatgc gtcaactggc 240 gcggctcagg ccgggcaggg ctgatgaacc tctacatcct caacatggcc atcgcggacc 300 tgggcattgt cctgtctctg cccgtgtgga tgctggaggt cacgctggac tacacctggc 360 totggggcag ottotoctgo ogottoacto actacttota otttgtcaac atgtatagca 420 gcatcttett cetgetgeee tteeetetea teacagtett caatgtgetg acageetgee 480 ggctgcggca gccaggacaa cccaagagcc ggcgccactg cctgctgctg tgcgcctacg 540 tggccgtctt tgtcatgtgc tggctgccct atcatgtgac cctgctgctg ctcacactgc 600 atgggaccca catctccctc cactgccacc tggtccacct gctctacttc ttctatgatg 660 teattgactg ettetecatg etgeactgtg teateaacce cateetttae aactttetea 720 gcccacactt ccggggccgg ctcctgaatg ctgtagtcca ttaccttcct aaggaccaga 780 ccaagggcgg gcacatgcgc ctcctcttcc tcctgttcca cccagcattc catcatcatc 840 accaaggtga tagccagcct gctgcagcag cccccaccc tgagccaagc ctgagctttc 900 aggcacacca tttgcttcca aatacttccc ccatctctcc cactcagcct cttacaccca 960 965 gctga <210> 68 <211> 320 <212> PRT <213> Homo sapiens <400> 68

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10

15

Ala Val Pro Thr Ser Asp Leu Gly Glu Ile His Asn Trp Thr Glu Leu Leu Asp Leu Phe Asn His Thr Leu Ser Glu Cys His Val Glu Leu Ser Gln Ser Thr Lys Arg Val Val Leu Phe Ala Leu Tyr Leu Ala Met Phe Val Val Gly Leu Val Glu Asn Leu Leu Val Ile Cys Val Asn Trp Arg Gly Ser Gly Arg Ala Gly Leu Met Asn Leu Tyr Ile Leu Asn Met Ala Ile Ala Asp Leu Gly Ile Val Leu Ser Leu Pro Val Trp Met Leu Glu Val Thr Leu Asp Tyr Thr Trp Leu Trp Gly Ser Phe Ser Cys Arg Phe Thr His Tyr Phe Tyr Phe Val Asn Met Tyr Ser Ser Ile Phe Phe Leu Leu Pro Phe Pro Leu Ile Thr Val Phe Asn Val Leu Thr Ala Cys Arg Leu Arg Gln Pro Gly Gln Pro Lys Ser Arg Arg His Cys Leu Leu Leu Cys Ala Tyr Val Ala Val Phe Val Met Cys Trp Leu Pro Tyr His Val Thr Leu Leu Leu Thr Leu His Gly Thr His Ile Ser Leu His Cys His Leu Val His Leu Leu Tyr Phe Phe Tyr Asp Val Ile Asp Cys Phe Ser Met Leu His Cys Val Ile Asn Pro Ile Leu Tyr Asn Phe Leu Ser Pro His Phe Arg Gly Arg Leu Leu Asn Ala Val Val His Tyr Leu Pro Lys Asp Gln Thr Lys Gly Gly His Met Arg Leu Leu Phe Leu Leu Phe

His Pro Ala Phe His His His Gln Gly Asp Ser Gln Pro Ala Ala 275 280 285

Ala Ala Pro His Pro Glu Pro Ser Leu Ser Phe Gln Ala His His Leu 290 295 300

Leu Pro Asn Thr Ser Pro Ile Ser Pro Thr Gln Pro Leu Thr Pro Ser 305 310 315 320

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<211> 549

<212> DNA

<213> Homo sapiens

<400> 69

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<211> 170

<212> PRT

<213> Homo sapiens

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Pro Ile Thr His Asn Pro Thr Asn Val Thr Leu Asn Lys Phe Ile Glu 20 25 30

Glu Leu Lys Lys Tyr Gly Ala Thr Thr Ile Val Arg Val Cys Glu Ala 35 40 45

Thr Tyr Asp Thr Thr Leu Val Glu Lys Glu Gly Ile His Val Leu Asn 50 55 60

Trp Pro Phe Gly Asp Gly Ala Pro Pro Ser Asn Gln Ile Val Ala Asp 65 70 75 80

Trp Leu His Phe Val Lys Ile Lys Phe Cys Glu Glu Pro Gly Cys Tyr 85 90 95

Ile Ala Val Asn Cys Ile Val Gly Leu Gly Lys Ala Pro Val Leu Val
100 105 110

Ala Leu Ala Ser Val Glu Gly Gly Met Lys His Glu Asp Ala Val Gln
115 120 125

Phe Ile Gly Gln Lys Arg Ser Gly Ala Phe Lys Ser Lys Gln Leu Leu 130 135 140

Tyr Leu Glu Lys Tyr His Pro Lys Met Arg Leu Arg Phe Lys Asp Ser 145 150 155 160

Asn Ser His Ile Asn Asn Cys Cys Ile Gln 165 170

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<213> Homo sapiens

<400> 71

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<211> 176

<212> PRT

<213> Homo sapiens

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Thr Tyr Lys Asn Met Arg Phe Pro Ile Thr His Asn Pro Thr Asn Val 20 25 30

Thr Leu Asn Lys Phe Ile Glu Glu Leu Lys Lys Tyr Gly Ala Thr Thr 35 40 45

Ile Val Arg Val Cys Glu Ala Thr Tyr Asp Thr Thr Leu Val Glu Lys 50 55 60

Glu Gly Ile His Val Leu Asn Trp Pro Phe Gly Asp Gly Ala Pro Pro 65 70 75 80

Ser Asn Gln Ile Val Ala Asp Trp Leu His Phe Val Lys Ile Lys Phe 85 90 95

Cys Glu Glu Pro Gly Cys Tyr Ile Ala Val Asn Cys Ile Val Gly Leu 100 105 110

Gly Lys Ala Pro Val Leu Val Ala Leu Ala Ser Val Glu Gly Gly Met 115 120 125

Lys His Glu Asp Ala Val Gln Phe Ile Gly Gln Lys Arg Ser Gly Ala 130 135 140

Phe Lys Ser Lys Gln Leu Leu Tyr Leu Glu Lys Tyr His Pro Lys Met 145 150 155 160

Arg Leu Arg Phe Lys Asp Ser Asn Ser Ala Ala Leu Gln Arg Phe Gln 165 170 175

<210> 73

<211> 1144

<212> DNA

<213> Homo sapiens

<400> 73

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<210> 74

<211> 355

<212> PRT

<213> Homo sapiens

<400> 74

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Met Glu Met Gly Arg Arg Met Asp Ala Pro Thr Ser Ala Ala Val Thr
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Arg Ala Phe Leu Glu Arg Gly His Thr Glu Ile Asp Thr Ala Phe Leu
35 40 45

Tyr Ser Asp Gly Gln Ser Glu Thr Ile Leu Gly Gly Leu Gly Leu Arg
50 55 60

Met Gly Ser Ser Asp Cys Arg Val Lys Ile Ala Thr Lys Ala Asn Pro 65 70 75 80

Trp Ile Gly Asn Ser Leu Lys Pro Asp Ser Val Arg Ser Gln Leu Glu 85 90 95

Thr Ser Leu Lys Arg Leu Gln Cys Pro Arg Val Asp Leu Phe Tyr Leu 100 105 110

| His | Ala | Pro 115 | Asp | His | Ser | Ala | Pro 120 | Val | Glu | Glu | Thr | Leu 125 | Arg | Ala | Cys |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Gln 130 | Leu | His | Gln | Glu | Gly 135 | Lys | Phe | Val | Glu | Leu 140 | Gly | Leu | Ser | Asn |
| Tyr 145 | Ala | Ala | Trp | Glu | Val 150 | Ala | Glu | Ile | Cys | Thr 155 | Leu | Cys | Lys | Ser | Asn 160 |
| Gly | Trp | Ile | Leu | Pro 165 | Thr | Val | Tyr | Gln | Gly 170 | Met | Tyr | Ser | Ala | Thr 175 | Thr |
| Arg | Gln | Val | Glu 180 | Thr | Glu | Leu | Phe | Pro 185 | Cys | Leu | Arg | His | Phe 190 | Gly | Leu |
| Arg | Phe | Tyr 195 | Ala | Tyr | Asn | Pro | Leu 200 | Ala | Asp | Gln | Ser | Pro 205 | Glu | Gly | Cys |
| Gly | Ser 210 | Phe | Trp | Gly | Thr | Leu 215 | Gly | Pro | Gly | Ala | Asp 220 | Cys | Cys | Leu | Pro |
| Ala 225 | Gly | Gly | Leu | Leu | Thr 230 | Gly | Lys | Tyr | Lys | Tyr 235 | Glu | Asp | Lys | Asp | Gly 240 |
| Lys | Gln | Pro | Val | Gly 245 | Arg | Phe | Phe | Gly | Thr 250 | Gln | Trp | Ala | Glu | Ile 255 | Tyr |
| Arg | Asn | Gln | Phe 260 | Trp | Lys | Glu | His | His 265 | Phe | Glu | Gly | Ile | Ala 270 | Leu | Val |
| Glu | Lys | Ala 275 | Leu | Gln | Ala | Ala | Tyr 280 | Gly | Ala | Ser | Ala | Pro 285 | Ser | Met | Thr |
| Ser | Ala 290 | Ala | Leu | Arg | Trp | Met 295 | | His | His | Ser | Gln 300 | Leu | Gln | Gly | Ala |
| His 305 | Gly | Asp | Ala | Val | Ile 310 | Leu | Gly | Met | Ser | Ser 315 | Leu | Glu | Gln | Leu | Glu 320 |
| Gln | Asn | Leu | Ala | Ala 325 | Ala | Glu | Glu | Gly | Pro 330 | Leu | Glu | Pro | Ala | Val 335 | Val |
| Asp | Ala | Phe | Asn 340 | Gln | Ala | Trp | His | Leu 345 | Phe | Ala | His | Glu | Cys 350 | Pro | Asn |
| Tyr | Phe | Ile 355 | | | | | | | | | | | | | |

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<213> Homo sapiens

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- Gln Arg Ser Gln Arg Ser Asp His Gln Arg Ser Gly Val Gly Gln Ala 35 40 45
- Pro Ser Pro Ile Ala Asn Thr Phe Leu His Tyr Arg Thr Ser Lys Val 50 55 60
- Arg Val Leu Arg Ala Ala Arg Leu Glu Arg Leu Val Gly Glu Leu Val
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- Phe Gly Asp Arg Glu Gln Asp Pro Ser Phe Met Pro Ala Phe Leu Ala 85 90 95
- Thr Tyr Arg Thr Phe Val Pro Thr Ala Cys Leu Leu Gly Phe Leu Leu 100 105 110
- Pro Pro Met Pro Pro Pro Pro Pro Gly Val Glu Ile Lys Lys Thr 115 120 125
- Ala Val Gln Asp Leu Ser Phe Asn Lys Asn Leu Arg Ala Val Val Ser 130 135 140
- Val Leu Gly Ser Trp Leu Gln Asp His Pro Gln Asp Phe Arg Asp Pro 145 150 155 160
- Pro Ala His Ser Asp Leu Gly Ser Val Arg Thr Phe Leu Gly Trp Ala 165 170 175
- Ala Pro Gly Ser Ala Glu Ala Gln Lys Ala Glu Lys Leu Leu Glu Asp 180 185 190
- Phe Leu Glu Glu Ala Glu Arg Glu Glu Glu Glu Pro Pro Gln Val 195 200 205
- Trp Ser Gly Pro Pro Arg Val Ala Gln Thr Ser Asp Pro Asp Ser Ser 210 215 220
- Glu Ala Cys Ala Glu Glu Glu Glu Gly Leu Met Pro Gln Gly Pro Gln 225 235 240

| Leu | Leu | Asp | Phe | Ser 245 | Val | Asp | Glu | Val | Ala 250 | Glu | Gln | Leu | Thr | Leu 255 | Ile |
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| Asp | Leu | Glu | Leu 260 | Phe | Ser | Lys | Val | Arg 265 | Leu | Tyr | Glu | Cys | Leu 270 | Gly | Ser |
| Val | Trp | Ser 275 | Gln | Arg | Asp | Arg | Pro 280 | Gly | Ala | Ala | Gly | Ala 285 | Ser | Pro | Thr |
| Val | Arg 290 | Ala | Thr | Val | Ala | Gln 295 | Phe | Asn | Thr | Val | Thr 300 | Gly | Cys | Val | Leu |
| Gly 305 | Ser | Val | Leu | Gly | Ala 310 | Pro | Gly | Leu | Ala | Ala 315 | Pro | Gln | Arg | Ala | Gln 320 |
| Arg | Leu | Glu | Lys | Trp 325 | Ile | Arg | Ile | Ala | Gln 330 | Arg | Cys | Arg | Glu | Leu 335 | Arg |
| Asn | Phe | Ser | Ser 340 | Leu | Arg | Ala | Ile | Leu 345 | Ser | Ala | Leu | Gln | Ser 350 | Asn | Pro |
| Ile | Tyr | Arg 355 | Leu | Lys | Arg | Ser | Trp 360 | Gly | Ala | Val | Ser | Arg 365 | Glu | Pro | Leu |
| Ser | Thr 370 | Phe | Arg | Lys | Leu | Ser 375 | Gln | Ile | Phe | Ser | Asp 380 | Glu | Asn | Asn | His |
| Leu 385 | Ser | Ser | Arg | Glu | Ile 390 | Leu | Phe | Gln | Glu | Glu 395 | Ala | Thr | Glu | Gly | Ser 400 |
| Gln | Glu | Glu | Asp | Asn 405 | Thr | Pro | Gly | Ser | Leu 410 | Pro | Ser | Lys | Pro | Pro 415 | Pro |
| Gly | Pro | Val | Pro 420 | Tyr | Leu | Gly | Thr | Phe 425 | Leu | Thr | Asp | Leu | Val 430 | Met | Leu |
| Asp | Thr | Ala 435 | Leu | Pro | Asp | Met | Leu 440 | Glu | Gly | Asp | Leu | Ile 445 | Asn | Phe | Glu |
| Lys | Arg 450 | Arg | Lys | Glu | Trp | Glu 455 | Ile | Leu | Ala | Arg | Ile 460 | Gln | Gln | Leu | Gln |
| Arg 465 | Arg | Cys | Gln | Ser | Tyr 470 | Thr | Leu | Ser | Pro | His 475 | Pro | Pro | Ile | Leu | Ala 480 |
| Ala | Leu | His | Ala | Gln 485 | Asn | Gln | Leu | Thr | Glu 490 | Glu | Gln | Ser | Tyr | Arg 495 | Leu |

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Arg Glu Lys Ser Ser Ser Pro Ser Gly Ser Pro Gly Asp Pro Ser Ser 530 540

Pro Thr Ser Ser Val Ser Pro Gly Ser Pro Pro Ser Ser Pro Arg Ser 545 550 555 560

Arg Asp Ala Pro Ala Gly Ser Pro Pro Ala Ser Pro Gly Pro Gln Gly 565 570 575

Pro Ser Thr Lys Leu Pro Leu Ser Leu Asp Leu Pro Ser Pro Arg Ser 580 585 590

Pro Val Thr Leu Asp Pro Phe Ser Ala Arg Val Pro Leu Pro Ala Gln 595 600 605

Gln Ser Ser Glu Ala Arg Val Ile Arg Val Ser Ile Asp Asn Asp His 610 615 620

Gly Asn Leu Tyr Arg Ser Ile Leu Leu Thr Ser Gln Asp Lys Ala Pro 625 630 635 640

Ser Val Val Arg Arg Ala Leu Gln Lys His Asn Val Pro Gln Pro Trp 645 650 655

Ala Cys Asp Tyr Gln Leu Phe Gln Val Leu Pro Gly Asp Arg Leu Leu 660 665 670

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Val Ser Pro Ser 705

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717

135

130

Ser Pro Ser Thr Thr Ser Lys Thr Val Thr Thr Ser Gly Thr Thr Asn

140

Asn Thr Val Thr Pro Thr Ser Gln Pro Val Arg Lys Ser Thr Phe Asp 145 150 155 160 Ala Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Gly Val Gln Ala 165 170 Val Ile Phe Phe Leu Tyr Lys Phe Cys Lys Ser Lys Glu Arg Asn Tyr 180 185 190 His Thr Leu 195 <210> 79 <211> 2082 <212> DNA <213> Homo sapiens <400> 79 cagcttcccc atggatcact ctccaaatag attctttaca cacaggtaat gtcactcagc 60 cetttgggte caaccettg teececagee eegagtggt getetteggg ggeeeteate 120 cattggcaag tgactgtcta ttcacatctc tcttcctgtt gttgagtgag tgagggaggg 180 agcctgccgg ggatccacag ctcccagttt ccactcactc attacacagt gctcttggcc 240 ctgcatgtgc tgtcacggcc atttggggtc tatatcctgt ctcttagagg acagggacta 300 ctctgtttca aatgccaggg ctacttatgg actcctattc aacctgcaaa accctacttg 420 aatgeteect cagttetgaa geeteectgg etgeteette cageeteece acaacaacaa 480 cagcaccacc actatataat ggctaaatct gttgagcagt tgccatgggc cagacactgt 540 gctgagtaca tggatatgtt ttcttcttta atcctcacaa cccctcgagt cagccccaag 600 ctaggctacc ctttggcaaa ttcacatcat tattcaatca agagcctctg gggagaaaag 660 ttggaaaacc cagccctcta cctggacaca gtccagagcc tatggattcc tgaagagccc 720 cctgtaccta caggaggcag cgtgagaatt aaaaaggacc ctgaacttgt ggtgaccgac 780 ctgcgttttg ggacgatacc cgtgaggctg ttccagccga aggcagcatc ctccagaccc 840 cggcgaggca tcatcttcta ccatggaggg gccacagtat ttgggagcct ggattgttac 900 catggcctgt gcaattatct ggcccgggag actgaatctg tacttctgat gattgggtac 960 cgcaagette etgaceacea tteecetgee etttteeaag actgcatgaa tgceteeatt 1020 cactteetga aggeeetgga aacetatggg gtggaceeet eeagggttgt ggtetgtgga 1080 cttccccqqa tccqqqctca qqttctqatt tatccaqttq tccaqqcatt ctqtttqcaq 1200 tcgccatcct ttcagcagaa ccaaaatgtc ccattacttt cccggaagtt catggtgact 1260 tetetqtqta actatetqqc cattqacete teetqqcqtq acqccatett qaacqqcact 1320 tgcgtacccc cagacgtctg gaggaagtac gagaagtggc tcacccctga caacatcccc 1380 aagaaattta agaacacagg ctaccaaccc tggtctcccg gcccttttaa tgaagctgcc 1440 tatctagaag ccaaacatat gctggatgta gaaaattcac ccctgatagc agatgatgag 1500

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<213> Homo sapiens

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Pro Lys Leu Gly Tyr Pro Leu Ala Asn Ser His His Tyr Ser Ile Lys 35 40 45

Ser Leu Trp Gly Glu Lys Leu Glu Asn Pro Ala Leu Tyr Leu Asp Thr 50 55 60

Val Gln Ser Leu Trp Ile Pro Glu Glu Pro Pro Val Pro Thr Gly Gly 65 70 75 80

Ser Val Arg Ile Lys Lys Asp Pro Glu Leu Val Val Thr Asp Leu Arg 85 90 95

Phe Gly Thr Ile Pro Val Arg Leu Phe Gln Pro Lys Ala Ala Ser Ser 100 105 110

Arg Pro Arg Arg Gly Ile Ile Phe Tyr His Gly Gly Ala Thr Val Phe 115 120 125

Gly Ser Leu Asp Cys Tyr His Gly Leu Cys Asn Tyr Leu Ala Arg Glu 130 135 140

Thr Glu Ser Val Leu Leu Met Ile Gly Tyr Arg Lys Leu Pro Asp His 145 150 155 160

His Ser Pro Ala Leu Phe Gln Asp Cys Met Asn Ala Ser Ile His Phe 165 170 175

Leu Lys Ala Leu Glu Thr Tyr Gly Val Asp Pro Ser Arg Val Val Cys Gly Glu Ser Val Gly Gly Ala Ala Val Ala Ala Ile Thr Gln Ala Leu Val Gly Arg Ser Asp Leu Pro Arg Ile Arg Ala Gln Val Leu Ile Tyr Pro Val Val Gln Ala Phe Cys Leu Gln Ser Pro Ser Phe Gln Gln Asn Gln Asn Val Pro Leu Leu Ser Arg Lys Phe Met Val Thr Ser Leu Cys Asn Tyr Leu Ala Ile Asp Leu Ser Trp Arg Asp Ala Ile Leu Asn Gly Thr Cys Val Pro Pro Asp Val Trp Arg Lys Tyr Glu Lys Trp Leu Thr Pro Asp Asn Ile Pro Lys Lys Phe Lys Asn Thr Gly Tyr Gln Pro Trp Ser Pro Gly Pro Phe Asn Glu Ala Ala Tyr Leu Glu Ala Lys His Met Leu Asp Val Glu Asn Ser Pro Leu Ile Ala Asp Asp Glu Val Ile Ala Gln Leu Pro Glu Ala Phe Leu Val Ser Cys Glu Asn Asp Ile Leu Arg Asp Asp Ser Leu Leu Tyr Lys Lys Arg Leu Glu Asp Gln Gly Val Arg Val Thr Trp Tyr His Leu Tyr Asp Gly Phe His Gly Ser Ile Ile Phe Phe Asp Lys Lys Ala Leu Ser Phe Pro Cys Ser Leu Lys Ile Val Asn Ala Val Val Ser Tyr Ile Lys Gly Ile

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<213> Homo sapiens

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<211> 307

<212> PRT

<213> Homo sapiens

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Leu Ala Ile Ile Ser Phe Ile Cys Leu Asp Ser Arg Leu His Ser Pro 35 40 45

Met Tyr Phe Phe Leu Cys Asn Phe Ser Leu Met Glu Met Val Val Thr 50 55 60

Ser Thr Val Val His Arg Met Leu Ala Asp Leu Leu Ser Thr His Lys 65 70 75 80

Thr Met Ser Leu Ala Lys Cys Leu Thr Gln Ser Phe Phe Tyr Phe Ser 85 90 95

Leu Gly Ser Ala Asn Phe Leu Ile Leu Met Val Met Ala Phe Asp Arg

100 105 110

Tyr Val Ala Ile Cys His Pro Leu Arg Tyr Pro Thr Ile Thr Asn Gly
115 120 125

Pro Val Cys Val Lys Leu Val Val Ala Cys Trp Val Val Gly Phe Leu 130 135 140

Ser Ile Val Ser Pro Thr Leu Gln Lys Thr Arg Leu Trp Phe Cys Gly 145 150 155 160

Pro Asn Ile Ile Gly His Tyr Phe Cys Asp Ser Ala Pro Leu Lys 165 170 175

Leu Ala Cys Ser Asp Thr Arg His Ile Glu Arg Met Asp Leu Phe Leu 180 185 190

Ser Leu Leu Phe Val Leu Thr Thr Met Leu Leu Ile Ile Leu Ser Tyr 195 200 205

Ile Leu Ile Val Ala Ala Val Leu His Ile Pro Ser Ser Ser Gly Cys 210 215 220

Gln Lys Ala Phe Ser Thr Cys Ala Pro His Leu Thr Val Val Leu 225 230 235 240

Gly Tyr Gly Ser Ala Ile Phe Ile Tyr Val Arg Pro Gly Lys Gly His 245 250 255

Ser Thr Tyr Leu Asn Lys Ala Val Ala Met Val Thr Ala Met Val Thr 260 265 270

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Ala Cys Arg 305

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<212> DNA

<213> Homo sapiens

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<210> 84
<211> 528
<212> PRT
<213> Homo sapiens
<400> 84
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|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Trp | Gly | Ser 35 | Leu | Leu | Ile | Tle | Leu 40 | Lys | Asn | Glu | Gly | Phe 45 | Tyr | Ser | Ser |
| Thr | Cys 50 | Pro | Ala | Glu | Ser | Ser 55 | Thr | Asn | Thr | Thr | Gln 60 | Asp | Glu | Gln | Arg |
| Arg 65 | Trp | Pro | Gly | Cys | Asp 70 | Gln | Gln | Asp | Glu | Met 75 | Leu | Asn | Leu | Gly | Phe 80 |
| Thr | Ile | Gly | Ser | Phe 85 | Val | Leu | Ser | Ala | Thr 90 | Thr | Leu | Pro | Leu | Gly 95 | Ile |
| Leu | Met | Asp | Arg 100 | Phe | Gly | Pro | Arg | Pro 105 | Val | Arg | Leu | Val | Gly 110 | Ser | Ala |
| Cys | Phe | Thr 115 | Ala | Ser | Cys | Thr | Leu 120 | Met | Ala | Leu | Ala | Ser 125 | Arg | Asp | Val |
| Glu | Ala 130 | Leu | Ser | Pro | Leu | Ile 135 | Phe | Leu | Ala | Leu | Ser 140 | Leu | Asn | Gly | Phe |
| Gly 145 | Gly | Ile | Cys | Leu | Thr 150 | Phe | Thr | Ser | Leu | Lys 155 | Leu | Ile | Tyr | Asp | Ala 160 |
| Gly | Val | Ala | Phe | Val 165 | Val | Ile | Met | Phe | Thr 170 | Trp | Ser | Gly | Leu | Ala 175 | Cys |
| Leu | Ile | Phe | Leu 180 | Asn | Cys | Thr | Leu | Asn 185 | Trp | Pro | Ile | Glu | Ala 190 | Phe | Pro |
| Ala | Pro | Glu 195 | Glu | Val | Asn | Tyr | Thr 200 | Lys | Lys | Ile | Lys | Leu 205 | Ser | Gly | Leu |
| Ala | Leu 210 | Asp | His | Lys | Val | Thr 215 | Gly | Asp | Leu | Phe | Tyr 220 | Thr | His | Val | Thr |
| Thr 225 | Met | Gly | Gln | Arg | Leu 230 | Ser | Gln | Lys | Ala | Pro 235 | Ser | Leu | Glu | Asp | Gly 240 |
| Ser | Asp | Ala | Phe | Met 245 | Ser | Pro | Gln | Asp | Val 250 | Arg | Gly | Thr | Ser | Glu 255 | Asn |
| Leu | Pro | Glu | Arg 260 | Ser | Val | Pro | Leu | Arg 265 | Lys | Ser | Leu | Cys | Ser 270 | Pro | Thr |

| Phe | Leu | Trp 275 | Ser | Leu | Leu | Thr | Met 280 | Cys | Met | Thr | Gln | Leu 285 | Arg | Ile | Ile |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Phe | Tyr 290 | Met | Ala | Ala | Val | Asn 295 | Lys | Met | Leu | Glu | Tyr 300 | Leu | Val | Thr | Gly |
| Gly 305 | Gln | Glu | His | Glu | Thr 310 | Asn | Glu | Gln | Gln | Gln 315 | Lys | Val | Ala | Glu | Thr 320 |
| Val | Gly | Phe | Tyr | Ser 325 | Ser | Val | Phe | Gly | Ala 330 | Met | Gln | Leu | Leu | Cys 335 | Leu |
| Leu | Thr | Cys | Pro 340 | Leu | Ile | Gly | Tyr | Ile 345 | Met | Asp | Trp | Arg | Ile 350 | Lys | Asp |
| Cys | Val | Asp 355 | Ala | Pro | Thr | Gln | Gly 360 | Thr | Val | Leu | Gly | Asp 365 | Ala | Arg | Asp |
| Gly | Val 370 | Ala | Thr | Lys | Ser | Ile 375 | Arg | Pro | Arg | Tyr | Cys 380 | Lys | Ile | Gln | Lys |
| Leu 385 | Thr | Asn | Ala | Ile | Ser 390 | Ala | Phe | Thr | Leu | Thr 395 | Asn | Leu | Leu | Leu | Val 400 |
| Gly | Phe | Gly | Ile | Thr 405 | Cys | Leu | Ile | Asn | Asn 410 | Leu | His | Leu | Gln | Phe 415 | Val |
| Thr | Phe | Val | Leu 420 | His | Thr | Ile | Val | Arg 425 | Gly | Phe | Phe | His | Ser 430 | Ala | Cys |
| Gly | Ser | Leu 435 | Tyr | Ala | Ala | Val | Phe 440 | Pro | Ser | Asn | His | Phe 445 | Gly | Thr | Leu |
| Thr | Gly 450 | Leu | Gln | Ser | Leu | Ile 455 | Ser | Ala | Val | Phe | Ala 460 | Leu | Leu | Gln | Gln |
| Pro 465 | Leu | Phe | Met | Ala | Met 470 | Val | Gly | Pro | Leu | Lys 475 | Gly | Glu | Pro | Phe | Trp 480 |
| Val | Asn | Leu | Gly | Leu 485 | Leu | Leu | Phe | Ser | Leu 490 | Leu | Gly | Phe | Leu | Leu 495 | Pro |
| Ser | Tyr | Leu | Phe 500 | Tyr | Tyr | Arg | Ala | Arg 505 | Leu | Gln | Gln | Glu | Tyr 510 | Ala | Ala |
| Asn | Gly | Met 515 | Gly | Pro | Leu | Lys | Val 520 | Leu | Ser | Gly | Ser | Glu 525 | Val | Thr | Ala |

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<212> DNA

<213> Homo sapiens

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<211> 99

<212> PRT

<213> Homo sapiens

<400> 86

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Ala Pro Glu Glu Leu Gly Cys Phe Val Gly Thr Ala Glu Ala Leu Arg
35 40 45

Cys Gln Glu Glu Asn Tyr Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys
50 55 60

Ala Cys Gly Ser Gly Gly Arg Cys Ala Val Leu Gly Leu Cys Cys Ser 65 70 75 80

Pro Asp Gly Cys His Ala Asp Pro Ala Cys Asp Ala Glu Ala Thr Phe 85 90 95

Ser Gln Arg

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tccaaggaat ggatcgaaca ggagaagcaa gcaggcttcg taatgaggcg tgcatcacca 180
atatgcacta agggcgaata a
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<211> 50
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<213> Homo sapiens
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                                 25
Lys Gln Ala Gly Phe Val Met Arg Arg Ala Ser Pro Ile Cys Thr Lys
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                             40
                                                  45
Gly Glu
     50
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tccaaggaat ggatcgaaca ggagaagcaa gcaggcttcg taatgaggcg tgcatcgcca 180
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<213> Homo sapiens

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Ser Pro Ile Cys Thr Lys Gly Glu 50 55

20

35

25

Glu Trp Ile Glu Gln Glu Lys Gln Ala Gly Phe Val Met Arg Arg Ala

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<210> 94 <211> 140 <212> PRT <213> Homo sapiens

<400> 94

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Ser Glu Asp Tyr Met Lys Glu Leu Gly Ile Gly Arg Ala Ser Arg Lys 20 25 30

Leu Gly Arg Leu Ala Lys Pro Thr Val Thr Ile Ser Thr Asp Gly Asp 35 40 45

Val Ile Thr Ile Lys Thr Lys Ser Ile Phe Lys Asn Asn Glu Ile Ser 50 55 60

Phe Lys Leu Gly Glu Glu Phe Glu Glu Ile Thr Pro Gly Gly His Lys
65 70 75 80

Thr Lys Ser Lys Val Thr Leu Asp Lys Glu Ser Leu Ile Gln Val Gln 85 90 95

Asp Trp Asp Gly Lys Glu Thr Thr Ile Thr Arg Lys Leu Val Asp Gly
100 105 110

Lys Met Val Val Glu Ser Thr Val Asn Ser Val Ile Cys Thr Arg Thr 115 120 125

Tyr Glu Lys Val Ser Ser Asn Ser Val Ser Asn Ser

140

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Met Val Glu Pro Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 10

408

Phe Glu Asp Tyr Met Lys Glu Leu Gly Phe Ala Ala Arg Asn Met Ala 25 20

Gly Leu Val Lys Pro Thr Val Thr Ile Ser Val Asp Gly Lys Met Met 35 40

Thr Ile Arg Thr Glu Ser Ser Phe Gln Asp Thr Lys Ile Ser Phe Lys 55

Leu Gly Glu Glu Phe Asp Glu Thr Thr Ala Asp Asn Arg Lys Val Lys 70 75

Ser Thr Ile Thr Leu Glu Asn Gly Ser Met Ile His Val Gln Lys Trp 85 90 95

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459

Gln Lys Trp Leu Gly Lys Glu Thr Thr Ile Lys Arg Lys Ile Val Asp 100 105

Lys Val Lys Ser Thr Ile Thr Leu Glu Asn Gly Ser Met Ile His Val

85

90

Glu Lys Met Val Val Glu Cys Lys Met Asn Asn Ile Val Ser Thr Arg

Ile Tyr Glu Lys Val
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<211> 1238

<212> DNA

<213> Homo sapiens

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<211> 411

<212> PRT

<213> Homo sapiens

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Glu His Leu Asp His Gln Ala Ala His Gln Pro Phe Pro Arg Pro Arg
20 25 30

Phe Arg Gln Glu Thr Gly His Pro Ser Leu Gln Arg Asp Phe Pro Arg

| Ser | Phe | Leu | Leu | Asp | Leu | Pro | Asn | Phe | Pro | Asp | Leu | Ser | Lys | Ala | Asp |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | 50 | | | | | 55 | | | | | 60 | | | | |

45

40

35

- Ile Asn Gly Gln Asn Pro Asn Ile Gln Val Thr Ile Glu Val Val Asp 65 70 75 80
- Gly Pro Asp Ser Glu Ala Asp Lys Asp Gln His Pro Glu Asn Lys Pro 85 90 95
- Ser Trp Ser Val Pro Ser Pro Asp Trp Arg Ala Trp Trp Gln Arg Ser 100 105 110
- Leu Ser Leu Ala Arg Ala Asn Ser Gly Asp Gln Asp Tyr Lys Tyr Asp 115 120 125
- Ser Thr Ser Asp Asp Ser Asn Phe Leu Asn Pro Pro Arg Gly Trp Asp 130 135 140
- His Thr Ala Pro Gly His Arg Thr Phe Glu Thr Lys Asp Gln Pro Glu 145 150 155 160
- Tyr Asp Ser Thr Asp Gly Glu Gly Asp Trp Ser Leu Trp Ser Val Cys 165 170 175
- Ser Val Thr Cys Gly Asn Gly Asn Gln Lys Arg Thr Arg Ser Cys Gly 180 185 190
- Tyr Ala Cys Thr Ala Thr Glu Ser Arg Thr Cys Asp Arg Pro Asn Cys 195 200 205
- Pro Gly Ile Glu Asp Thr Phe Arg Thr Ala Ala Thr Glu Val Ser Leu 210 215 220
- Leu Ala Gly Ser Glu Glu Phe Asn Ala Thr Lys Leu Phe Glu Val Asp 225 230 235 240
- Thr Asp Ser Cys Glu Arg Trp Met Ser Cys Lys Ser Glu Phe Leu Lys 245 250 255
- Lys Tyr Met His Lys Val Met Asn Asp Leu Pro Ser Cys Pro Cys Ser 260 265 270
- Tyr Pro Thr Glu Val Ala Tyr Ser Thr Ala Asp Ile Phe Asp Arg Ile 275 280 285
- Lys Arg Lys Asp Phe Arg Trp Lys Asp Ala Ser Gly Pro Lys Glu Lys

Leu Glu Ile Tyr Lys Pro Thr Ala Arg Tyr Cys Ile Arg Ser Met Leu 305 310 315 320

Ser Leu Glu Ser Thr Thr Leu Ala Ala Gln His Cys Cys Tyr Gly Asp $325 \hspace{1cm} 330 \hspace{1cm} 335$

Asn Met Gln Leu Ile Thr Arg Gly Lys Gly Ala Gly Thr Pro Asn Leu 340 345 350

Ile Gly Thr Glu Phe Ser Ala Glu Leu His Tyr Lys Val Asp Val Leu 355 360 365

Pro Trp Ile Ile Cys Lys Gly Asp Trp Ser Arg Tyr Asn Glu Ala Arg 370 375 380

Pro Pro Asn Asn Gly Gln Glu Cys Thr Glu Ser Pro Ser Asp Glu Asp 385 390 395 400

Tyr Ile Lys Gln Phe Gln Glu Ala Arg Glu Tyr
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<211> 1463

<212> DNA

<213> Homo sapiens

<400> 101

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<211> 454

<212> PRT

<213> Homo sapiens

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Phe Ser Thr Arg Asp Leu Leu Lys Glu Leu Thr Leu Gly Ala Ser Gln 20 25 30

Asp Glu Val Ala Ala Phe Phe Val Ala Asp Leu Gly Ala Ile Val Arg 35 40 45

Lys His Phe Cys Phe Leu Lys Cys Leu Pro Arg Val Arg Pro Phe Tyr 50 55 60

Ala Val Lys Cys Asn Ser Ser Pro Gly Val Leu Lys Val Leu Ala Gln 65 70 75 80

Leu Gly Leu Gly Phe Ser Cys Ala Asn Lys Ala Glu Met Glu Leu Val 85 90 95

Gln His Ile Gly Ile Pro Ala Ser Lys Ile Ile Cys Ala Asn Pro Cys 100 105 110

Lys Gln Ile Ala Gln Ile Lys Tyr Ala Ala Lys His Gly Ile Gln Leu 115 120 125

Leu Ser Phe Asp Asn Glu Met Glu Leu Ala Lys Val Val Lys Ser His 130 135 140

Pro Ser Ala Lys Met Val Leu Cys Ile Ala Thr Asp Asp Ser His Ser 145 150 155 160

Leu Ser Cys Leu Ser Leu Lys Phe Gly Val Ser Leu Lys Ser Cys Arg 165 170 175

| His | Leu | Leu | Glu 180 | Asn | Ala | Lys | Lys | His 185 | His | Val | Glu | Val | Val 190 | Gly | Val |
|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|--------------------------|-------------------|-------------------|--------------------------|--------------------------|--------------------------|--------------------------|-------------------|--------------------------|--------------------------|
| Ser | Phe | His 195 | Ile | Gly | Ser | Gly | Cys 200 | Pro | Asp | Pro | Gln | Ala 205 | Tyr | Ala | Gln |
| Ser | Ile 210 | Ala | Asp | Ala | Arg | Leu 215 | Val | Phe | Glu | Met | Gly 220 | Thr | Glu | Leu | Gly |
| His 225 | Lys | Met | His | Val | Leu 230 | Asp | Leu | Gly | Gly | Gly 235 | Phe | Pro | Gly | Thr | Glu 240 |
| Gly | Ala | Lys | Val | Arg 245 | Phe | Glu | Glu | Ile | Ala 250 | Ser | Val | Ile | Asn | Ser 255 | Ala |
| Leu | Asp | Leu | Tyr 260 | Phe | Pro | Glu | Gly | Cys 265 | Gly | Val | Asp | Ile | Phe 270 | Ala | Glu |
| Leu | Gly | Arg 275 | Tyr | Tyr | Val | Thr | Ser 280 | Ala | Phe | Thr | Val | Ala 285 | Val | Ser | Ile |
| Ile | Ala 290 | Lys | Lys | Glu | Val | Leu 295 | Leu | Asp | Gln | Pro | Gly 300 | Arg | Glu | Glu | Glu |
| Asn 305 | Gly | Ser | Thr | Ser | Lys 310 | Thr | Ile | Val | Tyr | His 315 | Leu | Asp | Glu | Gly | Val 320 |
| Tyr | Gly | Ile | Phe | Asn 325 | Ser | Val | Leu | Phe | _ | Asn | Ile | Cys | Pro | Thr | Pro |
| ~ 1 | | | | | | | | | 330 | | | | | 335 | |
| lle | Leu | Gln | Lys 340 | | Pro | Ser | Thr | Glu 345 | | Pro | Leu | Tyr | Ser 350 | | Ser |
| | | | 340 | Lys | | | | 345 | Gln | | | _ | 350 | Ser | Ser Leu |
| Leu | Trp | Gly 355 | 340 Pro | Lys Ala | Val | Asp | Gly 360 | 345 Cys | Gln Asp | Cys | Val | Ala 365 | 350 | Ser | Leu |
| Leu | Trp Leu 370 | Gly 355 Pro | 340 Pro Gln | Lys Ala Leu | Val His | Asp Val 375 | Gly 360 | 345 Cys Asp | Gln Asp Trp | Cys Leu | Val Val 380 | Ala 365 Phe | 350 Glu | Ser Gly Asn | Leu Met |
| Leu Trp Gly 385 | Trp Leu 370 Ala | Gly 355 Pro Tyr | 340 Pro Gln Thr | Lys Ala Leu Val | Val His Gly 390 | Asp Val 375 Met | Gly 360 Gly | 345 Cys Asp | Gln Asp Trp Pro | Cys Leu Phe 395 | Val Val 380 Trp | Ala 365 Phe Gly | 350 Glu Asp | Ser Gly Asn Gln | Leu Met Ala 400 |

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<400> 103

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<210> 104

<211> 402

<212> PRT

<213> Homo sapiens

<400> 104

| Met 1 | Ala | Gly | Tyr | Leu 5 | Ser | Glu | Ser | Asp | Phe 10 | Val | Met | Val | Glu | Glu 15 | Gly |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Phe | Ser | Thr | Arg 20 | Asp | Leu | Leu | Lys | Glu 25 | Leu | Thr | Leu | Gly | Ala 30 | Ser | Gln |
| Ala | Thr | Thr 35 | Ala | Glu | Met | Glu | Leu 40 | Val | Gln | His | Ile | Gly 45 | Ile | Pro | Ala |
| Ser | Lys 50 | Ile | Ile | Cys | Ala | Asn 55 | Pro | Cys | Lys | Gln | Ile 60 | Ala | Gln | Ile | Lys |
| Tyr 65 | Ala | Ala | Lys | His | Gly 70 | Ile | Gln | Leu | Leu | Ser 75 | Phe | Asp | Asn | Glu | Met 80 |
| Glu | Leu | Ala | Lys | Val 85 | Val | Lys | Ser | His | Pro 90 | Ser | Ala | Lys | Met | Val 95 | Leu |
| Cys | Ile | Ala | Thr 100 | Asp | Asp | Ser | His | Ser 105 | Leu | Ser | Cys | Leu | Ser 110 | Leu | Lys |
| Phe | Gly | Val 115 | Ser | Leu | Lys | Ser | Cys 120 | Arg | His | Leu | Leu | Glu 125 | Asn | Ala | Lys |
| Lys | His 130 | His | Val | Glu | Val | Val 135 | Gly | Val | Ser | Phe | His 140 | Ile | Gly | Ser | Gly |
| Cys 145 | Pro | Asp | Pro | Gln | Ala 150 | Tyr | Ala | Gln | Ser | Ile 155 | Ala | Asp | Ala | Arg | Leu 160 |
| Val | Phe | Glu | Met | Gly 165 | Thr | Glu | Leu | Gly | His 170 | Lys | Met | His | Val | Leu 175 | Asp |
| Leu | Gly | Gly | Gly 180 | Phe | Pro | Gly | Thr | Glu 185 | Gly | Ala | Lys | Val | Arg 190 | Phe | Glu |
| Glu | Ile | Ala 195 | Ser | Val | Ile | Asn | Ser 200 | Ala | Leu | Asp | Leu | Tyr 205 | Phe | Pro | Glu |
| Gly | Cys 210 | Gly | Val | Asp | Ile | Phe 215 | Ala | Glu | Leu | Gly | Arg 220 | Tyr | Tyr | Val | Thr |
| Ser 225 | Ala | Phe | Thr | Val | Ala 230 | Val | Ser | Ile | Ile | Ala 235 | Lys | Lys | Glu | Val | Leu 240 |
| Leu | Asp | Gln | Pro | Gly 245 | Arg | Glu | Glu | Glu | Asn 250 | Gly | Ser | Thr | Ser | Lys 255 | Thr |

Ile Val Tyr His Leu Asp Glu Gly Val Tyr Gly Ile Phe Asn Ser Val Leu Phe Asp Asn Ile Cys Pro Thr Pro Ile Leu Gln Lys Lys Pro Ser Thr Glu Gln Pro Leu Tyr Ser Ser Leu Trp Gly Pro Ala Val Asp Gly Cys Asp Cys Val Ala Glu Gly Leu Trp Leu Pro Gln Leu His Val Gly Asp Trp Leu Val Phe Asp Asn Met Gly Ala Tyr Thr Val Gly Met Gly Ser Pro Phe Trp Gly Thr Gln Ala Cys His Ile Thr Tyr Ala Met Ser Arg Val Ala Trp Glu Ala Leu Arg Arg Gln Leu Met Ala Ala Glu Gln Glu Asp Asp Val Glu Gly Val Cys Lys Pro Leu Ser Cys Gly Trp Glu Ile Thr Asp Thr Leu Cys Val Gly Pro Val Phe Thr Pro Ala Ser Ile Met

<210> 105 <211> 679 <212> DNA <213> Homo sapiens

<400> 105

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<210> 106

<211> 218

<212> PRT

<213> Homo sapiens

<400> 106

Met Ala Gly Tyr Leu Ser Glu Ser Asp Phe Val Met Val Glu Glu Gly 1 5 10 15

Phe Ser Thr Arg Asp Leu Leu Lys Glu Leu Thr Leu Gly Ala Ser Gln
20 25 30

Ala Thr Thr Asp Glu Val Ala Ala Phe Phe Val Ala Asp Leu Gly Ala 35 40 45

Ile Val Arg Lys His Phe Cys Phe Leu Lys Cys Leu Pro Arg Val Arg 50 55 60

Pro Phe Tyr Ala Val Lys Cys Asn Ser Ser Pro Gly Val Leu Lys Val 65 70 75 80

Leu Ala Gln Leu Gly Leu Gly Phe Ser Cys Ala Asn Ile Cys Pro Thr 85 90 95

Pro Ile Leu Gln Lys Lys Pro Ser Thr Glu Gln Pro Leu Tyr Ser Ser 100 · 105 110

Ser Leu Trp Gly Pro Ala Val Asp Gly Cys Asp Cys Val Ala Glu Gly
115 120 125

Leu Trp Leu Pro Gln Leu His Val Gly Asp Trp Leu Val Phe Asp Asn 130 135 140

Met Gly Ala Tyr Thr Val Gly Met Gly Ser Pro Phe Trp Gly Thr Gln 145 150 155 160

Ala Cys His Ile Thr Tyr Ala Met Ser Arg Val Ala Trp Glu Ala Leu 165 170 175

Arg Arg Gln Leu Met Ala Ala Glu Gln Glu Asp Asp Val Glu Gly Val 180 185 190

Cys Lys Pro Leu Ser Cys Gly Trp Glu Ile Thr Asp Thr Leu Cys Val 195 200 205

Gly Pro Val Phe Thr Pro Ala Ser Ile Met 210 215

<210> 107 <211> 2972 <212> DNA <213> Homo sapiens

<400> 107

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<211> 760

<212> PRT

<213> Homo sapiens

<400> 108

Met Leu Gln Ile Thr Glu Trp Arg Phe Leu Ala Arg Asp Glu Gly Glu
1 5 10 15

Ser Ala Val Ala Glu Asp Pro Thr Trp Gly Glu Asp Glu Glu Pro Ser 20 25 30

Ala Cys Thr Thr Asp Ser Trp Ala Gln Gly Ser Val Pro Val Leu His $35 \hspace{1cm} 40 \hspace{1cm} 45$

Ala Ser Thr Ser Glu Gly Leu Glu Asn Phe Gln Gly Glu Val His Ser 50 55 60

Ser Gly Ala Ser Pro Asp Ser Ser Ala Ile Ala Pro Ala Leu Pro Phe 65 70 75 80

Pro Thr Ser His Cys Pro Ser Ala Phe Pro Gln Asp Pro Gly Gly Val 85 90 95

Asp Arg Ile Pro Leu Gly Arg Ser Trp Met Gly Arg Gly Ser Gln Glu
100 105 110

Gln Met Glu Ser Trp Glu Pro Ser Pro Gln Leu Arg Val Thr Ser Ala 115 120 125

Pro Pro Pro Thr Ser Glu Leu Phe Gln Glu Ala Gly Pro Gly Gly Pro
130 135 140

Val Glu Glu Ala Asp Gly Gln Ser Arg Gly Leu Ser Ser Ala Gly Ser

| 145 | 150 | | 155 | 160 |
|------------------------|--------------------|--------------------|-----------------------|------------------|
| Leu Ser Ala Ser | Phe Gln Leu 165 | Ser Val Glu 170 | Glu Ala Pro Al | a Asp Asp 175 |
| Ala Asp Pro Ser 180 | Leu Asp Pro | Tyr Leu Val 185 | Ala Ser Pro Gl | |
| Thr Gly Arg Gly 195 | His Pro Leu | Gly Phe His 200 | Leu Ser Leu Gl 205 | u Asp Leu |
| Tyr Cys Cys Met 210 | Pro Gln Leu 215 | | Gly Asp Arg Le 220 | u Glu Leu |
| Arg Ser Glu Gly 225 | Val Pro Cys 230 | Ile Ala Ser | Gly Val Leu Va 235 | l Ser Tyr 240 |
| Pro Ser Val Gly | Gly Ala Thr 245 | Arg Pro Ser 250 | Ala Ser Cys Gl | n Gln Gln 255 |
| Arg Ala Gly His 260 | Ser Asp Val | Arg Leu Ser 265 | Ala His His Hi 27 | |
| Arg Arg Lys Ala 275 | Ala Val Lys | Arg Leu Asp 280 | Pro Ala Arg Le 285 | u Pro Cys |
| His Trp Val Arg 290 | Pro Leu Ala 295 | | Val Pro Asp Se 300 | r Gln Thr |
| Arg Pro Leu Glu 305 | Ala Tyr Arg 310 | Gly Arg Gln | Arg Gly Glu Ly 315 | s Thr Lys 320 |
| Ala Arg Ala Glu | Pro Gln Ala 325 | Leu Gly Pro 330 | Gly Thr Arg Va | l Ser Pro 335 |
| Ala Ala Phe Phe 340 | Pro Leu Arg | Pro Gly Ile 345 | Pro Phe Arg As | |
| Ser Gly Pro Ala 355 | Leu Leu Phe | Pro Thr Leu 360 | Asn Leu Gly Le 365 | u Ser Ser |
| Pro Ser Leu Glu 370 | Ser Lys Leu 375 | | Asn Ser Arg Il 380 | e Arg Phe |
| Leu Thr Thr His 385 | Pro Val Leu 390 | Pro Asp Val | Ala Arg Ser Ar | g Ser Pro 400 |

Lys Leu Trp Pro Ser Val Arg Trp Pro Ser Gly Trp Glu Gly Lys Ala

| Glu | Leu | Leu | Gly 420 | Glu | Leu | Trp | Ala | Gly 425 | Arg | Thr | Arg | Val | Pro 430 | Pro | Glr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly | Leu | Glu 435 | Leu | Ala | Asp | Arg | Glu 440 | Gly | Gln | Asp | Pro | Gly 445 | Arg | Trp | Pro |
| Arg | Thr 450 | Thr | Pro | Pro | Val | Leu 455 | Glu | Ala | Thr | Ser | Gln 460 | Val | Met | Trp | Lys |
| Pro 465 | Val | Leu | Leu | Pro | Glu 470 | Ala | Leu | Lys | Leu | Ala 475 | Pro | Gly | Val | Ser | Met 480 |
| Trp | Asn | Arg | Ser | Thr 485 | Gln | Val | Leu | Leu | Ser 490 | Ser | Gly | Val | Pro | Glu 495 | Gln |
| Glu | Asp | Lys | Glu 500 | Gly | Ser | Thr | Phe | Pro 505 | Pro | Val | Glu | Gln | His 510 | Pro | Ile |
| Gln | Thr | Gly 515 | Ala | Pro | Lys | Pro | Ser 520 | Ile | Ser | Pro | Ala | Gly 525 | Pro | Gly | Ser |
| Phe | Cys 530 | Tyr | Val | Ala | Val | Gly 535 | Cys | Thr | Gln | His | Pro 540 | Gly | Leu | Gly | Arg |
| Trp 545 | Leu | Cys | Leu | Pro | Tyr 550 | Ser | Gly | Leu | Leu | Gln 555 | Leu | His | Val | Gln | Leu 560 |
| Trp | Gln | Lys | Ser | His 565 | Pro | Trp | Asp | Leu | Gln 570 | Cys | Cys | Ser | Thr | Asp 575 | Leu |
| Thr | Gly | Lys | Ile 580 | Ala | Ile | Val | Thr | Gly 585 | Ala | Asn | Ser | Gly | Ile 590 | Gly | Lys |
| Val | Val | Ser 595 | Gln | Asp | Leu | Ala | Arg 600 | Cys | Gly | Ala | Gln | Val 605 | Ile | Leu | Thr |
| Cys | Gln 610 | Ser | Arg | Glu | Cys | Gly 615 | Gln | Gln | Ala | Leu | Ala 620 | Glu | Ile | Gln | Ala |
| Ala 625 | Ser | Asn | Ser | Asn | Arg 630 | Leu | Leu | Leu | Gly | Glu 635 | Val | Asp | Leu | Ser | Ser 640 |
| Met | Thr | Ser | Ile | Arg 645 | Ser | Phe | Ala | Arg | Arg 650 | Leu | Leu | Gln | Glu | Asn 655 | Pro |

Glu Ile His Leu Leu Val Asn Asn Ala Gly Val Ser Gly Phe Arg Arg

His Leu Pro Gln Gly Ala Trp Ile Ser Pro Leu Ser Leu Thr Met Leu 675 680 685

Gly Pro Phe Cys Ser Gln Ile Tyr Ser Lys Asp Leu Lys Gln Gly Val 690 695 700

Leu Pro Val Leu Tyr Leu Ser Leu Ala Glu Glu Pro Gly Gly Ile Ser 705 710 715 720

Gly Lys Tyr Phe Ser Ser Ser Cys Val Ile Thr Leu Pro Val Lys Ala 725 730 735

Ser Arg Asp Pro His Val Ala Gln Ser Leu Trp Asn Ala Ser Val Arg 740 745 750

Leu Thr Ser Leu Val Lys Met Asp 755 760

<210> 109

<211> 2077

<212> DNA

<213> Homo sapiens

<400> 109

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<210> 110

<211> 659

<212> PRT

<213> Homo sapiens

<400> 110

Met Glu Arg Trp Arg Asp Arg Leu Ala Leu Val Thr Gly Ala Ser Gly
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Gly Ile Gly Ala Ala Val Ala Arg Ala Leu Val Gln Gln Gly Leu Lys 20 25 30

Val Val Gly Cys Ala Arg Thr Val Gly Asn Ile Glu Glu Leu Ala Ala
35 40 45

Glu Cys Lys Ser Ala Gly Tyr Pro Gly Thr Leu Ile Pro Tyr Arg Cys
50 55 60

Asp Leu Ser Asn Glu Glu Asp Ile Leu Ser Met Phe Ser Ala Ile Arg 65 70 75 80

Ser Gln His Ser Gly Val Asp Ile Cys Ile Asn Asn Ala Gly Leu Ala 85 90 95

Arg Pro Asp Thr Leu Leu Ser Gly Ser Thr Ser Gly Trp Lys Asp Met
100 105 110

Phe Asn Val Asn Val Leu Ala Leu Ser Ile Cys Thr Arg Glu Ala Tyr 115 120 125

Gln Ser Met Lys Glu Arg Asn Val Asp Asp Gly His Ile Ile Asn Ile 130 135 140

| Asn 145 | Ser | Met | Ser | Gly | His 150 | Arg | Val | Leu | Pro | Leu 155 | Ser | Val | Thr | His | Phe 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Tyr | Ser | Ala | Thr | Lys 165 | Tyr | Ala | Val | Thr | Ala 170 | Leu | Thr | Glu | Gly | Leu 175 | Arg |
| Gln | Glu | Leu | Arg 180 | Glu | Ala | Gln | Thr | His 185 | Ile | Arg | Ala | Thr | Trp 190 | Gln | Leu |
| Arg | Arg | Glu 195 | Glu | Ala | Ala | Ala | Gly 200 | Tyr | Gln | Ala | Ala | Ile 205 | Thr | Val | Lys |
| Leu | Gly 210 | Phe | Cys | Gly | Leu | His 215 | Pro | Leu | Pro | Ser | Thr 220 | Ser | Pro | Arg | Pro |
| Gly 225 | Lys | Ala | Gln | Pro | Leu 230 | Arg | Arg | Pro | Ser | Leu 235 | Leu | Ala | Gln | Cys | Ile 240 |
| Ser | Pro | Gly | Val | Val 245 | Glu | Thr | Gln | Phe | Ala 250 | Phe | Lys | Leu | His | Asp 255 | Lys |
| Asp | Pro | Glu | Lys 260 | Ala | Ala | Ala | Thr | Tyr 265 | Glu | Gln | Met | Lys | Cys 270 | Leu | Lys |
| Pro | Glu | Asp 275 | Val | Ala | Glu | Ala | Val 280 | Ile | Tyr | Val | Leu | Ser 285 | Thr | Pro | Ala |
| His | Ile 290 | Gln | Ile | Gly | Asp | Ile 295 | Gln | Met | Arg | Pro | Thr 300 | Glu | Gln | Arg | Ala |
| Arg 305 | Arg | Arg | Arg | Leu | Ser 310 | Ser | Thr | Leu | His | Leu 315 | Gly | Val | Gly | Ser | Leu 320 |
| Gly | Ala | Asn | Cys | Gly 325 | Ala | Gly | Tyr | Arg | Ser 330 | Arg | Gly | Arg | Ser | Lys 335 | Gly |
| His | Arg | Val | Pro 340 | Gly | Gly | Ser | Cys | Ala 345 | Met | Ala | Leu | Leu | Ser 350 | Thr | Val |
| Arg | Gly | Ala 355 | Thr | Trp | Gly | Arg | Leu 360 | Val | Thr | Arg | His | Phe 365 | Ser | His | Ala |
| Ala | Arg 370 | His | Gly | Glu | Arg | Pro 375 | Gly | Gly | Glu | Glu | Leu 380 | Ser | Arg | Leu | Leu |
| Leu 385 | Asp | Asp | Leu | Val | Pro 390 | Thr | Ser | Arg | Leu | Glu 395 | Leu | Leu | Phe | Gly | Met 400 |

Thr Pro Cys Leu Leu Ala Leu Gln Ala Ala Arg Arg Ser Val Ala Arg Leu Leu Gln Ala Gly Lys Ala Gly Leu Gln Gly Lys Arg Ala Glu Leu Leu Arg Met Ala Glu Ala Arg Asp Ile Pro Val Leu Arg Pro Arg Arg Gln Lys Leu Asp Thr Met Cys Arg Tyr Gln Val His Gln Gly Val Cys Met Glu Val Ser Pro Leu Arg Pro Arg Pro Trp Arg Glu Ala Gly Glu Ala Ser Pro Gly Asp Asp Pro Gln Gln Leu Trp Leu Val Leu Asp Gly Ile Gln Asp Pro Arg Asn Phe Gly Ala Val Leu Arg Ser Ala His Phe Leu Gly Val Asp Lys Thr Lys Ala Gln Gln Gly Trp Leu Val Ala Gly Thr Val Gly Cys Pro Ser Thr Glu Asp Pro Gln Ser Ser Glu Ile Pro Ile Met Ser Cys Leu Glu Phe Leu Trp Glu Arg Pro Thr Leu Leu Val Leu Gly Asn Glu Gly Ser Gly Leu Ser Gln Glu Val Gln Ala Ser Cys Gln Leu Leu Thr Ile Leu Pro Arg Arg Gln Leu Pro Pro Gly Leu Glu Ser Leu Asn Val Ser Val Ala Ala Gly Ile Leu Leu His Ser Ile Cys Ser Gln Arg Lys Gly Phe Pro Thr Glu Gly Glu Arg Arg Gln Leu Leu Gln Asp Pro Gln Glu Pro Ser Ala Arg Ser Glu Gly Leu Ser Met Ala Gln His Pro Gly Leu Ser Ser Gly Pro Glu Lys Glu Arg Gln

<210> 111 <211> 3010 <212> DNA <213> Homo sapiens <400> 111 aatcttttt tttttttt ttttcgtaga taaaagtgca ttttatttcc ctagattgca 60 tttatttaat tcatataaca tgagaaactc ctccagtagc gtcaactagg gttgataaga 120 ataatcgata aagcaaaata aaaacacctt ctccaagatt ttgtaactgc aagcgaacgc 180 atggtggcgc tgttgactaa gaaggcgaat taaaccacag gcattgtgca tgctcggtga 240 cgcacggatc cagtgtggta aaccagcggt tgagagccca ggcagatttt tgagccagca 300 agtctgagcc tctggaaagg cttattcact aggccgtcta caaaggttgt ggggcaaaag 360 actgtttccc agetetgtet gaggttcage ttggegaeat teeetggaag agegtgaegg 420 aaagtgcaat ggaggcggga ggagagcgat ttcttagaca aaggcaagtc ttgcttctct 480 ttgtttttct gggagggtct ctggctgggt ccgagtcaag acgctattct gtggctgagg 540 aaaaagagaa gggcttttta atagccaacc tagcaaagga tctgggacta agggtagagg 600 aactggccgc gaggggggcc caagttgtgt ccaaagggaa caaacagcat tttcagctca 660 gtcatcagac aggtgatttg ctcctgaatg agaaattgga ccgggaggag ctatgcggcc 720 ccacagaacc atgcatacta cattttcaga tattactgca aaaccctttg caattcgtta 780 caaacgagct ccgtatcata gatgtaaatg accattctcc ggtattcttt gaaaatgaaa 840 tgcatctgaa aatcctagaa agcactctgc caggaacagt aattcctttg ggaaatgctg 900 aggacttgga tgtgggaaga aacagcctcc aaaactacac tatcactccg aattcccact 960 tccacgtacc cactcgcagt cgtagggacg gaaggaagta cccggaacta gtactgaaca 1020 gagccctgga tcgcgaggag cagcctgaga tcaggttaac cctcacagcg ctagatggcg 1080 ggagtccacc caggtccggc acggccctgg tacggattga agttgtggac atcaatgaca 1140 acgtcccaga gtttgcaaag ctgctctatg aggtgcagat cccggaggac agccccgttg 1200 gatcccaggt tgccatcgtc tctgccaggg atttagacat tggaactaat ggagaaatat 1260 cttatgcatt ttcccaagca tctgaagaca ttcgcaaaac gtttcgatta agtgcaaaat 1320 cgggagaact gcttttaaga cagaaactgg atttcgaatc catccagaca tacacagtaa 1380 atattcaggc gacagatggt gggggcctat ccggaaagtc tacagtcata gtccaggtgg 1440 ttgatgtcaa cgacaaccca ccggaactga ccttgtcttc agtaaacagc cctattcctg 1500 agaactcggg agagactgta ctggctgttt tcagtgtttc tgatctagac tctggagaca 1560 acggaagagt gatgtgttcc attgagaaca atctcccctt cttcctgaaa ccatctgtag 1620 agaattttta caccctagtg tcagaaggcg cgctggacag agagaccaga tccgagtaca 1680 acattaccat cactatcact gacctgggga cacccaggct gaaaaccaag tacaacataa 1740 cegtgetggt cteegaegte aatgacaacg ceeeegeett caeecaaate teetacaeee 1800 tgttcgtccg cgagaacaac agccccgccc tgcacatcgg cagtgtcagc gccacagaca 1860 gagactcagg caccaacgcc caggtaacct actcgctgct gccgccccag gacccgcacc 1920 tgcccctctc ttccctggtc tccatcaacg cggacaacgg ccacctgttt gccctcaggt 1980 cgctggacta cgaggccctg caggcgttcg agttccgcgt gggcgccaca gaccgtggct 2040 ecceggettt gageagegag gegetggtge gegtgetggt getggaegee aacgaeaact 2100

cgcccttcgt gctgtacccg ctgcagaacg gctccgcgcc ctgcaccgag ctggtgcccc 2160 gggcggctga gccgggctac ctggtgacca aggtggtggc ggtggacggc gactcgggcc 2220

agaacgcctg gctgtcgtac cagctgcta aggccacgga gcccgggctg ttcggcgtgt ggggcgcacaa tggcgaagtg cgcaccgca ggctgctgag ggagcgcac gctgccaagc 2340 agaggctggt ggtgctggtc aaggacaatg gcgagcctcc gcgctcggcc accgccacgc 2400 cggcccaggc ccaggccgac ttgctcacc agccctacct gctgctcccg gaggcggcac 2460 cggcccaggc ccaggccgac ttgctcaccg tctacctggt ggtggcgttg gcctcggtt 2520 cttcgctct cctcttctcg gtgctcctgt tcgtggcgt gcggctgtc aggaggagca 2580 gggcggccc ggtgggcgc tgctcggtc ccgagggcc ccttccagg cagatggtg ccgagggcc ctgccgggac ctgcccaga gctaccaga gctaccagta cgaggtgtg ctgactggag 2700 gctccggac aaatgagtc aagttccta agccaattat ccccaacttc gttgctcagg 2760 gtgcagaaga ggttagcgag gcaaatccca gtttcaggaa gagctttgaa ttcacttaag 2820 tgttaataaa gatctactga ggctagtctc gtttaatttg tggaaagtcc tttttact 2880 ctttgcccat tggaggtgc tcctttatt agaaagtac catcttattc caattctatg 2940 catgttactg gtattataa atgtatgagt ttttttgcgg tataataaat gtaaatttc 3000 tttgtattct

<210> 112

<211> 796

<212> PRT

<213> Homo sapiens

<400> 112

Met Glu Ala Gly Gly Glu Arg Phe Leu Arg Gln Arg Gln Val Leu Leu 1 5 10 15

Leu Phe Val Phe Leu Gly Gly Ser Leu Ala Gly Ser Glu Ser Arg Arg 20 25 30

Tyr Ser Val Ala Glu Glu Lys Glu Lys Gly Phe Leu Ile Ala Asn Leu 35 40 45

Ala Lys Asp Leu Gly Leu Arg Val Glu Glu Leu Ala Ala Arg Gly Ala 50 55 60

Gln Val Val Ser Lys Gly Asn Lys Gln His Phe Gln Leu Ser His Gln 65 70 75 80

Thr Gly Asp Leu Leu Leu Asn Glu Lys Leu Asp Arg Glu Glu Leu Cys
85 90 95

Gly Pro Thr Glu Pro Cys Ile Leu His Phe Gln Ile Leu Leu Gln Asn $100 \hspace{1.5cm} 105 \hspace{1.5cm} 110 \hspace{1.5cm}$

Pro Leu Gln Phe Val Thr Asn Glu Leu Arg Ile Ile Asp Val Asn Asp 115 120 125

His Ser Pro Val Phe Phe Glu Asn Glu Met His Leu Lys Ile Leu Glu 130 135 140

| Ser Thr Leu 145 | Pro Gly | Thr Val | Ile | Pro | Leu | Gly 155 | Asn | Ala | Glu | Asp | Leu 160 |
|--------------------|----------------|----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Asp Val Gly | Arg Asn 165 | Ser Leu | Gln | Asn | Tyr 170 | Thr | Ile | Thr | Pro | Asn 175 | Ser |
| His Phe His | Val Pro 180 | Thr Arg | Ser | Arg 185 | Arg | Asp | Gly | Arg | Lys 190 | Tyr | Pro |
| Glu Leu Val 195 | Leu Asn | Arg Ala | Leu 200 | Asp | Arg | Glu | Glu | Gln 205 | Pro | Glu | Ile |
| Arg Leu Thr 210 | Leu Thr | Ala Leu 215 | _ | Gly | Gly | Ser | Pro 220 | Pro | Arg | Ser | Gly |
| Thr Ala Leu 225 | Val Arg | Ile Glu 230 | Val | Val | Asp | Ile 235 | Asn | Asp | Asn | Val | Pro 240 |
| Glu Phe Ala | Lys Leu 245 | Leu Tyr | Glu | Val | Gln 250 | Ile | Pro | Glu | Asp | Ser 255 | Pro |
| Val Gly Ser | Gln Val 260 | Ala Ile | Val | Ser 265 | Ala | Arg | Asp | Leu | Asp 270 | Ile | Gly |
| Thr Asn Gly 275 | Glu Ile | Ser Tyr | Ala 280 | Phe | Ser | Gln | Ala | Ser 285 | Glu | Asp | Ile |
| Arg Lys Thr 290 | Phe Arg | Leu Ser 295 | | Lys | Ser | Gly | Glu 300 | Leu | Leu | Leu | Arg |
| Gln Lys Leu 305 | Asp Phe | Glu Ser 310 | Ile | Gln | Thr | Tyr 315 | Thr | Val | Asn | Ile | Gln 320 |
| Ala Thr Asp | Gly Gly 325 | Gly Leu | Ser | Gly | Lys 330 | Ser | Thr | Val | Ile | Val 335 | Gln |
| Val Val Asp | Val Asn 340 | Asp Asn | Pro | Pro 345 | Glu | Leu | Thr | Leu | Ser 350 | Ser | Val |
| Asn Ser Pro 355 | Ile Pro | Glu Asn | Ser 360 | Gly | Glu | Thr | Val | Leu 365 | Ala | Val | Phe |
| Ser Val Ser 370 | Asp Leu | Asp Ser 375 | | Asp | Asn | Gly | Arg 380 | Val | Met | Cys | Ser |
| Ile Glu Asn 385 | Asn Leu | Pro Phe | Phe | Leu | Lys | Pro 395 | Ser | Val | Glu | Asn | Phe 400 |

Tyr Thr Leu Val Ser Glu Gly Ala Leu Asp Arg Glu Thr Arg Ser Glu Tyr Asn Ile Thr Ile Thr Ile Thr Asp Leu Gly Thr Pro Arg Leu Lys Thr Lys Tyr Asn Ile Thr Val Leu Val Ser Asp Val Asn Asp Asn Ala Pro Ala Phe Thr Gln Ile Ser Tyr Thr Leu Phe Val Arg Glu Asn Asn Ser Pro Ala Leu His Ile Gly Ser Val Ser Ala Thr Asp Arg Asp Ser Gly Thr Asn Ala Gln Val Thr Tyr Ser Leu Leu Pro Pro Gln Asp Pro His Leu Pro Leu Ser Ser Leu Val Ser Ile Asn Ala Asp Asn Gly His Leu Phe Ala Leu Arg Ser Leu Asp Tyr Glu Ala Leu Gln Ala Phe Glu Phe Arg Val Gly Ala Thr Asp Arg Gly Ser Pro Ala Leu Ser Ser Glu Ala Leu Val Arg Val Leu Val Leu Asp Ala Asn Asp Asn Ser Pro Phe Val Leu Tyr Pro Leu Gln Asn Gly Ser Ala Pro Cys Thr Glu Leu Val Pro Arg Ala Ala Glu Pro Gly Tyr Leu Val Thr Lys Val Val Ala Val Asp Gly Asp Ser Gly Gln Asn Ala Trp Leu Ser Tyr Gln Leu Leu Lys Ala Thr Glu Pro Gly Leu Phe Gly Val Trp Ala His Asn Gly Glu Val Arg Thr Ala Arg Leu Leu Arg Glu Arg Asp Ala Ala Lys Gln Arg Leu Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg Ser Ala Thr Ala

Thr Leu His Val Leu Leu Val Asp Gly Phe Ser Gln Pro Tyr Leu Leu Leu Pro Glu Ala Ala Pro Ala Gln Ala Gln Ala Asp Leu Leu Thr Val Tyr Leu Val Val Ala Leu Ala Ser Val Ser Ser Leu Phe Leu Phe Ser Val Leu Leu Phe Val Ala Val Arg Leu Cys Arg Arg Ser Arg Ala Ala Ser Val Gly Arg Cys Ser Val Pro Glu Gly Pro Phe Pro Gly Gln Met Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu Val Cys Leu Thr Gly Gly Ser Gly Thr Asn Glu Phe Lys Phe Leu Lys Pro Ile Ile Pro Asn Phe Val Ala Gln Gly Ala Glu Arg Val Ser Glu Ala Asn Pro Ser Phe Arg Lys Ser Phe Glu Phe Thr <210> 113 <211> 261 <212> PRT <213> Homo sapiens <400> 113 Met Ile Tyr Lys Cys Pro Met Cys Arg Glu Phe Phe Ser Glu Arg Ala Asp Leu Phe Met His Gln Lys Val His Thr Ala Glu Lys Pro His Lys Cys Asp Lys Cys Asp Lys Gly Phe Phe His Ile Ser Glu Leu His Ile His Trp Arg Asp His Thr Gly Glu Lys Val Tyr Lys Cys Asp Asp Cys Gly Lys Asp Phe Ser Thr Thr Thr Lys Leu Asn Arg His Lys Lys Ile His Thr Val Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe \$85\$ 90 95

Asn Trp Ser Pro His Leu Gln Ile His Met Arg Val His Thr Gly Glu
100 105 110

Lys Pro Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser 115 120 125

Asn Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Phe Lys 130 135 140

Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu Cys Met 145 150 155 160

His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Tyr Glu Cys 165 170 175

Gly Lys Ala Phe Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val 180 185 190

His Thr Gly Glu Lys Pro Tyr Arg Cys Cys Gly Cys Gly Lys Ala Phe 195 200 205

Ser Gln Ser Ser Ser Leu Cys Ile His Gln Arg Val His Thr Gly Glu 210 215 220

Lys Pro Phe Lys Cys Asp Glu Cys Gly Lys Ala Phe Ser Gln Ser Thr 225 230 235 240

Ser Leu Cys Ile His Gln Arg Val His Thr Lys Glu Arg Asn His Leu 245 250 255

Lys Ile Ser Val Ile 260

<210> 114

<211> 184

<212> PRT

<213> Homo sapiens

<400> 114

Val His Thr Ala Glu Lys Pro His Lys Cys Asp Lys Cys Asp Lys Gly
1 5 10 15

Phe Phe His Ile Ser Glu Leu His Ile His Trp Arg Asp His Thr Gly Glu Lys Val Tyr Lys Cys Asp Asp Cys Gly Lys Asp Phe Ser Thr Thr Thr Lys Leu Asn Arg His Lys Lys Ile His Thr Val Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Asn Trp Ser Ser His Leu Gln Ile His Met Arg Val His Thr Gly Glu Glu Pro Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser Asn Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Phe Lys Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Ser Gln Arg Ser Ser Leu Cys Ile His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Arg Cys Cys Gly Cys Gly Lys Ala <210> 115 <211> 183 <212> PRT <213> Homo sapiens <400> 115 Val His Thr Ala Glu Lys Pro His Lys Cys Asp Lys Cys Asp Lys Gly Phe Phe His Ile Ser Glu Leu His Ile His Trp Arg Asp His Thr Gly Glu Lys Val Tyr Lys Cys Asp Asp Cys Gly Lys Asp Phe Ser Thr Thr

Thr Lys Leu Asn Arg His Lys Lys Ile His Thr Val Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Asn Trp Ser Ser His Leu Gln Ile His Met Arg Val His Thr Gly Glu Glu Pro Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser Asn Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Phe Lys Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu Cys Met His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Ser Gln Arg Ser Ser Leu Cys Ile His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Arg Cys Cys Gly Cys Gly Lys <210> 116 <211> 1147 <212> PRT <213> Homo sapiens <400> 116 Met Pro Val Lys Lys Gly Cys Gln Gly Pro Pro Lys Gly Met Leu Arg Pro Cys Val Pro Gly Phe Ser Val Cys Ala Ser Gln Ser Leu Ile Ser

Pro Ala Glu Val Pro Gly Leu Arg Trp Ala Cys Leu Gln Glu Gln Leu

Val Leu Gly Ser Gly Asn Ser Val Glu Leu Ser Cys His Pro Pro Gly

Arg Gly Pro Met Glu Leu Thr Val Gly Val Lys Gly Ser Ala Gly Leu

- Pro Gly Thr Ser Ser Trp Gly Ser Thr Ile Val Ala Pro Pro Gly Ser 85 90 95
- Gly Ile Pro Pro Leu Pro Pro Arg Arg Arg His Ser Thr Arg Ser Leu
 100 105 110
- Ala Cys Cys Asn Ser Ile His Ser Ser Gly Ala Ala Ser Thr Val Gln
 115 120 125
- Ala Gly Gly Arg Gly Gly Gln Gly Gln Arg Ala Ala Phe Pro Gly Gly
 130 135 140
- Arg Thr Leu Pro Ser Pro Val Thr Arg Lys Thr Val Thr Val His Pro 145 150 155 160
- Glu Ser His Cys Gln Gln Leu His Val Asn Ser Ser Pro Lys Asp Thr 165 170 175
- Arg Glu Thr Gln Ala Ser Gly Pro Met Gly Thr Leu Gly Val Arg Ala 180 185 190
- Leu Ala Arg Gln Thr Gly Ala Val Tyr Lys Ser Arg Gly Pro Pro Gln
 195 200 205
- Gln Val Asp Arg Lys Glu Gln Ile Lys Gly Lys Pro Tyr Glu Thr His 210 215 220
- Leu Gln Arg Asn Gln Pro Ile Gln Glu Lys Thr Arg Phe Arg Ala Pro 225 230 235 240
- Leu Ala His Pro Arg Gly Arg Pro Cys Arg Pro Val Leu Ala Gln Leu 245 250 255
- Lys His Pro Pro Pro Tyr Pro Ser Leu Leu Lys Gly Ala Leu Cys Thr 260 265 270
- Gly Ala Glu Arg Phe Leu Ser Lys Ala Leu Trp Leu Ser Leu Ser Ser 275 280 285
- Pro Ser Thr Leu His Pro Thr Leu Ser Cys Ser Lys Gly Pro Cys Leu 290 295 300
- Pro Glu Gln Asn Thr Pro Ser Pro Arg Leu Tyr Gly Ser Arg Ala Gln 305 310 315
- Leu Arg Pro Lys Val Val Lys Gly Pro Phe Arg Ser Pro Lys Cys Ala

| | 325 | 330 | | 335 |
|------------------------|--------------------|--------------------|--------------------|--------------------|
| Gly Gln Leu Thr | - | Lys Ser Leu 345 | Val Pro Cys | Gly His Arg 350 |
| Glu Ala Met Ile 355 | Ala Ala Cys | Pro His Gly 360 | Lys Ala Phe 365 | Trp Ser Leu |
| His Val Arg Val | Gln Leu Trp 375 | Gln Gln Arg | Thr Phe Pro 380 | Val Leu Glu |
| Ile Leu Ser Val 385 | Trp Gln Gly 390 | Leu Gly Thr | Pro Thr Gln 395 | Pro Pro Ser 400 |
| Ala Ala Ser Cys | Gln Leu Trp 405 | Glu Asp Val 410 | Asp Trp Cys | Leu Val His 415 |
| Leu Ser Ser Cys 420 | | Arg Ser Val 425 | Asp Lys Ala | Gln Val Ser 430 |
| Ser Lys Ala Thr 435 | Thr Glu Asn | Ala Gln Asp 440 | Val Ile Arg 445 | Ala Leu Lys |
| Met Pro Gly Arg 450 | Val Glu Gly 455 | Lys Met Gln | Lys Leu Gln 460 | Glu Gly Lys |
| Val Asn Leu Glu 465 | Lys Asp Leu 470 | Glu Lys Glu | Ser Asn Arg 475 | Asp Ala Val 480 |
| Thr Ala Leu Arg | Thr Val Asp | Asp Leu Val 490 | Ile Ile Lys | Pro Met His 495 |
| Leu Ser Gly His | | Ile His Leu 505 | His Leu Cys | Ser Ser Gln 510 |
| Glu Glu Ala Ile 515 | Arg Ala Ala | Gln Trp Leu 520 | Val Gln Glu 525 | Ala Leu Pro |
| Leu Val Pro Trp 530 | Gly Lys Asp 535 | Leu Gln Trp | Gln His Gly 540 | Thr Tyr Asn |
| Ala Leu Ser Ala 545 | Asp Asp Ala 550 | Val Gln Ser | Pro Pro Asp 555 | Cys Ser Glu 560 |
| Asp Ala Thr Asn | Ser Cys Leu 565 | Thr Ile Thr 570 | Arg Val Thr | Glu Cys Ile 575 |

Arg Glu Ser Leu Cys Phe Lys Gln Cys Leu Thr Gly Gln Phe Leu Pro

| 580 | | 585 | 590 |
|------------------------|------------------|--------------------------|----------------------------|
| Glu Gln Val His 595 | | Phe Ser Trp Ser (| Gln Ile Lys Asn Ser 605 |
| Ala His Gly Thr 610 | Phe Cys Lys 615 | | Ala Phe Ser Asp Val 620 |
| Val Ile Glu Phe | Ser Pro Glu 630 | Glu Trp Ala Cys 1 | Leu Asp Pro Ala Gln |
| 625 | | 635 | 640 |
| Arg Asn Leu Tyr | Arg Asp Val 1 | Met Phe Glu Asn 1 | Tyr Arg Asn Leu Val |
| | 645 | 650 | 655 |
| Ser Leu Asp Leu | | Gln Asp Met Lys i | Asp Leu Cys Gln Lys |
| 660 | | 665 | 670 |
| Val Thr Leu Thr | | Ser Trp Gly Leu <i>i</i> | Asp Asn Leu His Leu |
| 675 | | 680 | 685 |
| Val Lys Asp Trp 690 | Arg Thr Val . | | Gly Gln Lys Glu Tyr 700 |
| Cys Asn Arg Leu | Thr Gln Cys | Ser Ser Thr Lys 3 | Ser Lys Ile Phe Gln |
| 705 | 710 | 715 | 720 |
| Cys Ile Glu Cys | Gly Arg Asn | Phe Ser Trp Arg : | Ser Ile Leu Thr Glu |
| | 725 | 730 | 735 |
| His Lys Arg Ile | _ | Glu Lys Pro Tyr 1 | Lys Cys Glu Glu Cys |
| 740 | | 745 | 750 |
| Gly Lys Val Phe | | Ser Asn Leu Thr 1 | Lys His Lys Arg Ile |
| 755 | | 760 | 765 |
| His Thr Gly Glu 770 | Lys Pro Tyr 7775 | | Cys Gly Lys Val Phe 780 |
| Asn Trp Trp Ser | Gln Leu Thr . | Asn His Lys Lys : | Ile His Thr Gly Glu |
| 785 | 790 | 795 | 800 |
| Lys Pro Tyr Lys | Cys Asp Glu | Cys Asp Lys Val 1 | Phe Asn Trp Trp Ser |
| | 805 | 810 | 815 |
| Gln Leu Thr Ser | | Ile His Ser Gly (| Glu Lys Pro Tyr Pro |
| 820 | | 825 | 830 |
| Cys Glu Glu Cys | Gly Lys Ala | Phe Thr Gln Phe | Ser Asn Leu Thr Gln |

835

His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Lys Glu Cys 850 855 860

845

- Cys Lys Ala Phe Asn Lys Phe Ser Asn Leu Thr Gln His Lys Arg Ile 865 870 875 880
- His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys Gly Asn Val Phe 885 890 895
- Asn Glu Cys Ser His Leu Thr Arg His Arg Arg Ile His Thr Gly Glu 900 905 910
- Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys Ala Phe Thr Gln Phe Ala 915 920 925
- Ser Leu Thr Arg His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Gln 930 935 940
- Cys Glu Glu Cys Gly Lys Thr Phe Asn Arg Cys Ser His Leu Ser Ser 945 950 955 960
- His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Lys Cys Glu Glu Cys 965 970 975
- Gly Arg Thr Phe Thr Gln Phe Ser Asn Leu Thr Gln His Lys Arg Ile 980 985 990
- His Thr Gly Glu Lys Pro Tyr Lys Cys Lys Glu Cys Gly Lys Ala Phe 995 1000 1005
- Asn Lys Phe Ser Ser Leu Thr Gln His Arg Arg Ile His Thr Gly Val 1010 1015 1020
- Lys Pro Tyr Lys Cys Glu Glu Cys Gly Lys Val Phe Lys Gln Cys Ser 1025 1030 1035 1040
- His Leu Thr Ser His Lys Arg Ile His Thr Gly Glu Lys Pro Tyr Lys
 1045 1050 1055
- Cys Lys Glu Cys Gly Lys Ala Phe Tyr Gln Ser Ser Ile Leu Ser Lys 1060 1065 1070
- His Lys Arg Ile His Thr Glu Glu Lys Pro Tyr Lys Cys Glu Glu Cys 1075 1080 1085
- Gly Lys Ala Phe Asn Gln Phe Ser Ser Leu Thr Arg His Lys Arg Ile

1090 1095 1100

His Thr Gly Glu Lys Arg Tyr Lys Cys Lys Glu Cys Gly Lys Gly Phe 1105 1110 1115

Tyr Gln Ser Ser Ile His Ser Lys Tyr Lys Arg Ile Tyr Thr Gly Glu 1125 1130 1135

Glu Pro Asp Lys Cys Lys Lys Cys Gly Ser Leu 1140 1145

<210> 117

<211> 606

<212> PRT

<213> Homo sapiens

<400> 117

Met Ala Val Thr Phe Glu Asp Val Thr Ile Ile Phe Thr Trp Glu Glu
1 5 10 15

Trp Lys Phe Leu Asp Ser Ser Gln Lys Arg Leu Tyr Arg Glu Val Met 20 25 30

Trp Glu Asn Tyr Thr Asn Val Met Ser Val Glu Asn Trp Asn Glu Ser 35 40 45

Tyr Lys Ser Gln Glu Glu Lys Phe Arg Tyr Leu Glu Tyr Glu Asn Phe 50 55 60

Ser Tyr Trp Gln Gly Trp Trp Asn Ala Gly Ala Gln Met Tyr Glu Asn 65 70 75 80

Gln Asn Tyr Gly Glu Thr Val Gln Gly Thr Asp Ser Lys Asp Leu Thr 85 90 95

Gln Gln Asp Arg Ser Gln Cys Gln Glu Trp Leu Ile Leu Ser Thr Gln
100 105 110

Val Pro Gly Tyr Gly Asn Tyr Glu Leu Thr Phe Glu Ser Lys Ser Leu 115 120 125

Arg Asn Leu Lys Tyr Lys Asn Phe Met Pro Trp Gln Ser Leu Glu Thr 130 135 140

Lys Thr Thr Gln Asp Tyr Gly Arg Glu Ile Tyr Met Ser Gly Ser His 145 150 155 160

| Gly | Phe | Gln | Gly | Gly 165 | Arg | Tyr | Arg | Leu | Gly 170 | Ile | Ser | Arg | Lys | Asn 175 | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser | Met | Glu | Lys 180 | Glu | Gln | Lys | Leu | Ile 185 | Val | Gln | His | Ser | Tyr 190 | Ile | Pro |
| Val | Glu | Glu 195 | Ala | Leu | Pro | Gln | Tyr 200 | Val | Gly | Val | Ile | Cys 205 | Gln | Glu | Asp |
| Leu | Leu 210 | Arg | Asp | Ser | Met | Glu 215 | Glu | Lys | Tyr | Cys | Gly 220 | Cys | Asn | Lys | Cys |
| Lys 225 | Gly | Ile | Tyr | Tyr | Trp 230 | Asn | Ser | Arg | Cys | Val 235 | Phe | His | Lys | Arg | Asn 240 |
| Gln | Pro | Gly | Glu | Asn 245 | Leu | Cys | Gln | Cys | Ser 250 | Ile | Arg | Lys | Ala | Cys 255 | Phe |
| Ser | Gln | Arg | Ser 260 | Asp | Leu | Tyr | Arg | His 265 | Pro | Arg | Asn | His | Ile 270 | Gly | Lys |
| Lys | Leu | Tyr 275 | Gly | Cys | Asp | Glu | Val 280 | Asp | Gly | Asn | Phe | His 285 | Gln | Ser | Ser |
| Gly | Val 290 | His | Phe | His | Gln | Arg 295 | Val | His | Ile | Gly | Glu 300 | Val | Pro | Tyr | Ser |
| Cys 305 | Asn | Ala | Cys | Gly | Lys 310 | Ser | Phe | Ser | Gln | Ile 315 | Ser | Ser | Leu | His | Asn 320 |
| His | Gln | Arg | Val | His 325 | Thr | Glu | Glu | Lys | Phe 330 | Tyr | Lys | Ile | Glu | Cys 335 | Asp |
| Lys | Asp | Leu | Ser 340 | Arg | Asn | Ser | Leu | Leu 345 | His | Ile | His | Gln | Arg 350 | Leu | His |
| Ile | Gly | Glu 355 | Lys | Pro | Phe | Lys | Cys 360 | Asn | Gln | Cys | Gly | Lys 365 | Ser | Phe | Asn |
| Arg | Ser 370 | Ser | Val | Leu | His | Val 375 | His | Gln | Arg | Val | His 380 | Thr | Gly | Glu | Lys |
| Pro 385 | Tyr | Lys | Cys | Asp | Glu 390 | Cys | Gly | Lys | Gly | Phe 395 | Ser | Gln | Ser | Ser | Asn 400 |
| Leu | Arg | Ile | His | Gln 405 | Leu | Val | His | Thr | Gly 410 | Glu | Lys | Ser | Tyr | Lys 415 | Cys |

Glu Asp Cys Gly Lys Gly Phe Thr Gln Arg Ser Asn Leu Gln Ile His
420 425 430

Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys Asp Asp Cys Gly 435 440 445

Lys Asp Phe Ser His Ser Ser Asp Leu Arg Ile His Gln Arg Val His 450 455 460

Thr Gly Glu Lys Pro Tyr Thr Cys Pro Glu Cys Gly Lys Gly Phe Ser 465 470 475 480

Lys Ser Ser Lys Leu His Thr His Gln Arg Val His Thr Gly Glu Lys
485 490 495

Pro Tyr Lys Cys Glu Glu Cys Gly Lys Gly Phe Ser Gln Arg Ser His 500 505 510

Leu Leu Ile His Gln Arg Val His Thr Gly Glu Lys Pro Tyr Lys Cys 515 520 525

His Asp Cys Gly Lys Gly Phe Ser His Ser Ser Asn Leu His Ile His 530 540

Gln Arg Val His Thr Gly Glu Lys Pro Tyr Gln Cys Ala Lys Cys Gly 545 550 555 560

Lys Gly Phe Ser His Ser Ser Ala Leu Arg Ile His Gln Arg Val His 565 570 575

Ala Gly Glu Lys Pro Tyr Lys Cys Arg Glu Tyr Tyr Lys Gly Phe Asp 580 585 590

His Asn Ser His Leu His Asn Asn His Arg Arg Gly Asn Leu 595 600 605

<210> 118

<211> 24

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: zinc finger C2H2 consensus pattern sequence

<400> 118

Tyr Lys Cys Pro Phe Asp Cys Gly Lys Ser Phe Ser Arg Lys Ser Asn

Leu Lys Arg His Leu Arg Thr His 20 <210> 119 <211> 23 <212> PRT <213> Homo sapiens <400> 119 Tyr Lys Cys Pro Met Cys Arg Glu Phe Phe Ser Glu Arg Ala Asp Leu 10 15 Phe Met His Gln Lys Ile His 20 <210> 120 <211> 23 <212> PRT <213> Homo sapiens <400> 120 His Lys Cys Asp Lys Cys Asp Lys Gly Phe Phe His Ile Ser Glu Leu 5 His Ile His Trp Arg Asp His 20 <210> 121 <211> 23 <212> PRT <213> Homo sapiens <400> 121 Tyr Lys Cys Asp Asp Cys Gly Lys Asp Phe Ser Thr Thr Thr Lys Leu 5 10 15 Asn Arg His Lys Lys Ile His 20 <210> 122 <211> 23

10

15

1

<212> PRT

<213> Homo sapiens <400> 122 Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Asn Trp Ser Ser His Leu 5 10 15 Gln Ile His Met Arg Val His 20 <210> 123 <211> 23 <212> PRT <213> Homo sapiens <400> 123 Tyr Val Cys Ser Glu Cys Gly Arg Gly Phe Ser Asn Ser Ser Asn Leu 10 Cys Met His Gln Arg Val His 20 <210> 124 <211> 23 <212> PRT <213> Homo sapiens <400> 124 Phe Lys Cys Glu Glu Cys Gly Lys Ala Phe Arg His Thr Ser Ser Leu 1 5 10 15 Cys Met His Gln Arg Val His 20 <210> 125 <211> 23 <212> PRT <213> Homo sapiens <400> 125 Tyr Lys Cys Tyr Glu Cys Gly Lys Ala Phe Ser Gln Ser Ser Leu 5 10 15 Cys Ile His Gln Arg Val His

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<210> 126
<211> 23
<212> PRT
<213> Homo sapiens
<400> 126
Tyr Arg Cys Cys Gly Cys Gly Lys Ala Phe Ser Gln Ser Ser Gly Leu
                                     10
Cys Ile His Gln Arg Val His
             20
<210> 127
<211> 23
<212> PRT
<213> Homo sapiens
<400> 127
Phe Lys Cys Asp Glu Cys Gly Lys Ala Phe Ser Gln Ser Thr Ser Leu
 1
                  5
                                     10
                                                          15
Cys Ile His Gln Arg Val His
             20
<210> 128
<211> 388
<212> PRT
<213> Homo sapiens
<400> 128
Met Lys Trp Leu Leu Leu Gly Leu Val Ala Leu Ser Glu Cys Ile
                                     10
Met Tyr Lys Val Pro Leu Ile Arg Lys Lys Ser Leu Arg Arg Thr Leu
             20
                                 25
Ser Glu Arg Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Leu Asn
         35
                             40
Pro Ala Arg Lys Tyr Phe Pro Gln Trp Glu Ala Pro Thr Leu Val Asp
     50
                         55
Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile
```

75

70

| Gly | Ile | Gly | Thr | Pro 85 | Ala | Gln | Asp | Phe | Thr 90 | Val | Val | Phe | Asp | Thr 95 | Gly |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser | Ser | Asn | Leu 100 | Trp | Val | Pro | Ser | Val 105 | Tyr | Cys | Ser | Ser | Leu 110 | Ala | Cys |
| Thr | Asn | His 115 | Asn | Arg | Phe | Asn | Pro 120 | Glu | Asp | Ser | Ser | Thr 125 | Tyr | Gln | Ser |
| Thr | Ser 130 | Glu | Thr | Val | Ser | Ile 135 | Thr | Tyr | Gly | Thr | Gly 140 | Ser | Met | Thr | Gly |
| Ile 145 | Leu | Gly | Tyr | Asp | Thr 150 | Val | Gln | Val | Gly | Gly 155 | Ile | Ser | Asp | Thr | Asn 160 |
| Gln | Ile | Phe | Gly | Leu 165 | Ser | Glu | Thr | Glu | Pro 170 | Gly | Ser | Phe | Leu | Tyr 175 | Tyr |
| Ala | Pro | Phe | Asp 180 | Gly | Ile | Leu | Gly | Leu 185 | Ala | Tyr | Pro | Ser | Ile 190 | Ser | Ser |
| Ser | Gly | Ala 195 | Thr | Pro | Val | Phe | Asp 200 | Asn | Ile | Trp | Asn | Gln 205 | Gly | Leu | Val |
| Ser | Gln 210 | Asp | Leu | Phe | Ser | Val 215 | Tyr | Leu | Ser | Ala | Asp 220 | Asp | Gln | Ser | Gly |
| Ser 225 | Val | Val | Ile | Phe | Gly 230 | Gly | Ile | Asp | Ser | Ser 235 | Tyr | Tyr | Thr | Gly | Ser 240 |
| Leu | Asn | Trp | Val | Pro 245 | Val | Thr | Val | Glu | Gly 250 | Tyr | Trp | Gln | Ile | Thr 255 | Val |
| Asp | Ser | Ile | Thr 260 | Met | Asn | Gly | Glu | Ala 265 | Ile | Ala | Cys | Ala | Glu 270 | Gly | Cys |
| Gln | Ala | Ile 275 | Val | Asp | Thr | Gly | Thr 280 | Ser | Leu | Leu | Thr | Gly 285 | Pro | Thr | Ser |
| Pro | Ile 290 | Ala | Asn | Ile | Gln | Ser 295 | Asp | Ile | Gly | Ala | Ser 300 | Glu | Asn | Ser | Asp |
| Gly 305 | Asp | Met | Val | Val | Ser 310 | Cys | Ser | Ala | Ile | Ser 315 | Ser | Leu | Pro | Asp | Ile 320 |
| Val | Phe | Thr | Ile | Asn 325 | Gly | Val | Gln | Tyr | Pro 330 | Val | Pro | Pro | Ser | Ala 335 | Tyr |

.

Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn 340 345 350

Leu Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile 355 360 365

Arg Gln Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn Gln Val Gly Leu 370 380

Ala Pro Val Ala 385

<210> 129

<211> 388

<212> PRT

<213> Homo sapiens

<400> 129

Met Lys Trp Leu Leu Leu Gly Leu Val Ala Leu Ser Glu Cys Ile 1 5 10 15

Met Tyr Lys Val Pro Leu Ile Arg Lys Lys Ser Leu Arg Arg Thr Leu 20 25 30

Ser Glu Arg Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Leu Asn 35 40 45

Pro Ala Arg Lys Tyr Phe Pro Gln Trp Lys Ala Pro Thr Leu Val Asp 50 55 60

Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile 65 70 75 80

Gly Ile Gly Thr Pro Ala Gln Asp Phe Thr Val Leu Phe Asp Thr Gly 85 90 95

Ser Ser Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu Ala Cys 100 105 110

Thr Asn His Asn Arg Phe Asn Pro Glu Asp Ser Ser Thr Tyr Gln Ser 115 120 125

Thr Ser Glu Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met Thr Gly 130 135 140

Gln Ile Phe Gly Leu Ser Glu Thr Glu Pro Gly Ser Phe Leu Tyr Tyr Ala Pro Phe Asp Gly Ile Leu Gly Leu Ala Tyr Pro Ser Ile Ser Ser Ser Gly Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly Leu Val Ser Gln Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln Ser Gly Ser Val Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr Gly Ser Leu Asn Trp Val Pro Val Thr Val Glu Gly Tyr Trp Gln Ile Thr Val Asp Ser Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu Gly Cys Gln Ala Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Gly Pro Thr Ser Pro Ile Ala Asn Ile Gln Ser Asp Ile Gly Ala Ser Glu Asn Ser Asp Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn Leu Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile

Ala Pro Val Ala

<210> 130

Arg Gln Tyr Phe Thr Val Phe Glu Arg Ala Asn Asn Gln Val Gly Leu

```
<211> 388
<212> PRT
<213> Homo sapiens
<400> 130
Met Lys Trp Leu Lei
1
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Met Lys Trp Leu Leu Leu Gly Leu Val Ala Leu Ser Glu Cys Ile 1 5 10 15

Met Tyr Lys Val Pro Leu Ile Arg Lys Lys Ser Phe Arg Arg Thr Leu 20 25 30

Ser Glu Arg Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Leu Asn 35 40 45

Pro Ala Arg Lys Tyr Phe Pro Gln Trp Lys Ala Pro Thr Leu Val Asp 50 55 60

Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile 65 70 75 80

Gly Ile Gly Thr Pro Ala Gln Asp Phe Thr Val Leu Phe Asp Thr Gly 85 90 95

Ser Ser Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu Ala Cys
100 105 110

Thr Asn His Asn Arg Phe Asn Pro Glu Asp Ser Ser Thr Tyr Gln Ser 115 120 125

Thr Ser Glu Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met Thr Gly 130 135 140

Ile Leu Gly Tyr Asp Thr Val Gln Val Gly Gly Ile Ser Asp Thr Asn 145 150 155 160

Gln Ile Phe Gly Leu Ser Glu Thr Glu Pro Gly Ser Phe Leu Tyr Tyr 165 170 175

Ala Pro Phe Asp Gly Ile Leu Gly Leu Ala Tyr Pro Ser Ile Ser Ser 180 185 190

Ser Gly Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly Leu Val 195 200 205

Ser Gln Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln Ser Gly 210 215 220

Ser Val Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr Gly Ser

| 225 | 225 | | | | | | | | 235 | | | | | |
|---------|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|
| Leu Asn | Trp | Val | Pro 245 | Val | Thr | Val | Glu | Gly 250 | Tyr | Trp | Gln | Ile | Thr 255 | Val |

Asp Ser Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu Gly Cys

Gln Ala Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Gly Pro Thr Ser

Pro Ile Ala Asn Ile Gln Ser Asp Ile Gly Ala Ser Glu Asn Ser Asp

Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile

Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr

Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn

Leu Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile

Arg Gln Tyr Phe Thr Val Phe Glu Arg Ala Asn Asn Gln Val Gly Leu

Ala Pro Val Ala

<210> 131

<211> 388

<212> PRT

<213> Homo sapiens

<400> 131

Met Lys Trp Leu Leu Leu Gly Leu Val Ala Leu Ser Glu Cys Ile

Met Tyr Lys Val Pro Leu Ile Arg Lys Lys Ser Leu Arg Arg Thr Leu

Ser Glu Arg Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Leu Asn

| Pro | Ala 50 | Arg | Lys | Tyr | Phe | Pro 55 | Gln | Trp | Glu | Ala | Pro 60 | Thr | Leu | Val | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu 65 | Gln | Pro | Leu | Glu | Asn 70 | Tyr | Leu | Asp | Met | Glu 75 | Tyr | Phe | Gly | Thr | Ile 80 |
| Gly | Ile | Gly | Thr | Pro 85 | Ala | Gln | Asp | Phe | Thr 90 | Val | Leu | Phe | Asp | Thr 95 | Gly |
| Ser | Ser | Asn | Leu 100 | Trp | Val | Pro | Ser | Val 105 | Tyr | Cys | Ser | Ser | Leu 110 | Ala | Cys |
| Thr | Asn | His 115 | Asn | Arg | Phe | Asn | Pro 120 | Glu | Asp | Ser | Ser | Thr 125 | Tyr | Gln | Ser |
| Thr | Ser 130 | Glu | Thr | Val | Ser | Ile 135 | Thr | Tyr | Gly | Thr | Gly 140 | Ser | Met | Thr | Gly |
| Ile 145 | Leu | Gly | Tyr | Asp | Thr 150 | Val | Gln | Val | Gly | Gly 155 | Ile | Ser | Asp | Thr | Asn 160 |
| Gln | Ile | Phe | Gly | Leu 165 | Ser | Glu | Thr | Glu | Pro 170 | Gly | Ser | Phe | Leu | Tyr 175 | Tyr |
| Ala | Pro | Phe | Asp 180 | Gly | Ile | Leu | Gly | Leu 185 | Ala | Tyr | Pro | Ser | Ile 190 | Ser | Ser |
| Ser | Gly | Ala 195 | Thr | Pro | Val | Phe | Asp 200 | Asn | Ile | Trp | Asn | Gln 205 | Gly | Leu | Val |
| Ser | Gln 210 | Asp | Leu | Phe | Ser | Val 215 | Tyr | Leu | Ser | Ala | Asp 220 | Asp | Lys | Ser | Gly |
| Ser 225 | Val | Val | Ile | Phe | Gly 230 | Gly | Ile | Asp | Ser | Ser 235 | Tyr | Tyr | Thr | Gly | Ser 240 |
| Leu | Asn | Trp | Val | Pro 245 | Val | Thr | Val | Glu | Gly 250 | Tyr | Trp | Gln | Ile | Thr 255 | Val |
| Asp | Ser | Ile | Thr 260 | Met | Asn | Gly | Glu | Thr 265 | Ile | Ala | Cys | Ala | Glu 270 | Gly | Cys |
| Gln | Ala | Ile 275 | Val | Asp | Thr | Gly | Thr 280 | Ser | Leu | Leu | Thr | Gly 285 | Pro | Thr | Ser |
| Pro | Ile 290 | Ala | Asn | Ile | Gln | Ser 295 | Asp | Ile | Gly | Ala | Ser 300 | Glu | Asn | Ser | Asp |

Gly Asp Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile Val Phe Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr Ile Leu Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn Val Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile Arg Gln Tyr Phe Thr Val Phe Glu Arg Ala Asn Asn Gln Val Gly Leu Ala Pro Val Ala <210> 132 <211> 328 <212> PRT <213> Macaca fuscata <400> 132 Met Lys Trp Leu Leu Leu Gly Leu Val Ala Leu Ser Glu Cys Ile Ile His Lys Val Pro Leu Val Arg Lys Lys Ser Leu Arg Arg Asn Leu Ser Glu His Gly Leu Leu Lys Asp Phe Leu Lys Lys His Asn Phe Asn Pro Ala Ser Lys Tyr Phe Pro Gln Ala Glu Ala Pro Thr Leu Ile Asp Glu Gln Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile Gly Ile Gly Thr Pro Ala Gln Asp Phe Thr Val Ile Phe Asp Thr Gly Ser Ser Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu Ala Cys Thr Asn His Asn Arg Phe Asn Pro Gln Asp Ser Ser Thr Tyr Gln Ser

Thr Ser Gly Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met Thr Gly Ile Leu Gly Tyr Asp Thr Val Gln Val Gly Gly Ile Ser Asp Thr Asn Gln Ile Phe Gly Leu Ser Glu Thr Glu Pro Gly Ser Phe Leu Tyr Tyr Ala Pro Phe Asp Gly Ile Leu Gly Leu Ala Tyr Pro Ser Ile Ser Ser Ser Gly Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly Leu Val Ser Gln Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln Ser Gly Ser Val Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr Gly Ser Leu Asn Trp Val Pro Val Ser Val Glu Gly Tyr Trp Gln Ile Ser Val Asp Ser Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu Gly Cys Gln Ala Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Ile Ser Gly Phe Gln Gly Met Asp Val Pro Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile Arg Gln Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn

Gln Val Gly Leu Ala Pro Val Ala

<210> 133

<211> 369

<212> PRT

<213> Homo sapiens

<400> 133

Lys Val Pro Leu Ile Arg Lys Lys Ser Leu Arg Arg Thr Leu Ser Glu

| 1 . | 5 | 10 | 15 |
|-----|---|----|----|
| | | | |

| Arg | Gly | Leu | Leu | Lys | Asp | Phe | Leu | Lys | Lys | His | Asn | Leu | Asn | Pro | Ala |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | | 20 | | | | | 25 | | | | | 30 | | |

- Arg Lys Tyr Phe Pro Gln Trp Glu Ala Pro Thr Leu Val Asp Glu Gln
 35 40 45
- Pro Leu Glu Asn Tyr Leu Asp Met Glu Tyr Phe Gly Thr Ile Gly Ile 50 55 60
- Gly Thr Pro Ala Gln Asp Phe Thr Val Leu Phe Asp Thr Gly Ser Ser 65 70 75 80
- Asn Leu Trp Val Pro Ser Val Tyr Cys Ser Ser Leu Ala Cys Thr Asn 85 90 95
- His Asn Arg Phe Asn Pro Glu Asp Ser Ser Thr Tyr Gln Ala Thr Ser 100 105 110
- Glu Thr Val Ser Ile Thr Tyr Gly Thr Gly Ser Met Thr Gly Ile Leu 115 120 125
- Gly Tyr Asp Thr Val Gln Val Gly Gly Ile Ser Asp Thr Asn Gln Ile 130 135 140
- Phe Gly Leu Ser Glu Thr Glu Pro Gly Ser Phe Leu Tyr Tyr Ala Pro 145 150 155 160
- Phe Asp Gly Ile Leu Gly Leu Ala Tyr Pro Ser Ile Ser Ser Gly
 165 170 175
- Ala Thr Pro Val Phe Asp Asn Ile Trp Asn Gln Gly Leu Val Ser Gln
 180 185 190
- Asp Leu Phe Ser Val Tyr Leu Ser Ala Asp Asp Gln Ser Gly Ser Val
 195 200 205
- Val Ile Phe Gly Gly Ile Asp Ser Ser Tyr Tyr Thr Gly Ser Leu Asn 210 215 220
- Trp Val Pro Val Thr Val Glu Gly Tyr Trp Gln Ile Thr Val Asp Ser 225 230 235 240
- Ile Thr Met Asn Gly Glu Ala Ile Ala Cys Ala Glu Gly Cys Gln Ala
 245 250 255
- Ile Val Asp Thr Gly Thr Ser Leu Leu Thr Gly Pro Thr Ser Pro Ile

260 265 270

Ala Asn Ile Gln Ser Asp Ile Gly Ala Ser Glu Asn Ser Asp Gly Asp 275 280 285

Met Val Val Ser Cys Ser Ala Ile Ser Ser Leu Pro Asp Ile Val Phe 290 295 300

Thr Ile Asn Gly Val Gln Tyr Pro Val Pro Pro Ser Ala Tyr Ile Leu 305 310 315 .320

Gln Ser Glu Gly Ser Cys Ile Ser Gly Phe Gln Gly Met Asn Leu Pro $325 \hspace{1cm} 330 \hspace{1cm} 335$

Thr Glu Ser Gly Glu Leu Trp Ile Leu Gly Asp Val Phe Ile Arg Gln 340 345 350

Tyr Phe Thr Val Phe Asp Arg Ala Asn Asn Gln Val Ser Leu Ala Pro 355 360 365

Val

<210> 134

<211> 374

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Eukaryotic aspartyl protease domain sequence

<400> 134

Arg Ile Pro Leu Lys Lys Val Pro Ser Leu Arg Glu Lys Leu Ser Glu

1 5 10 15

Lys Gly Val Leu Leu Asp Phe Leu Val Lys Arg Lys Tyr Glu Pro Thr 20 25 30

Lys Lys Leu Thr Gly Gly Ala Ser Ser Ser Arg Ser Ala Val Glu Pro 35 40 45

Leu Leu Asn Tyr Leu Asp Ala Glu Tyr Tyr Gly Thr Ile Ser Ile Gly 50 55 60

Thr Pro Pro Gln Lys Phe Thr Val Val Phe Asp Thr Gly Ser Ser Asp 65 70 75 80

| Leu | Trp | Val | Pro | Ser 85 | Val | Tyr | Cys | Thr | Ser 90 | Ser | Tyr | Ala | Cys | Lys 95 | Gly |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Gly | Thr | Phe 100 | Asp | Pro | Ser | Lys | Ser 105 | Ser | Thr | Туқ | Lys | Asn 110 | Leu | Gly |
| Thr | Thr | Phe 115 | Ser | Ile | Ser | Tyr | Gly 120 | Asp | Gly | Ser | Ser | Ala 125 | Ser | Gly | Phe |
| Leu | Gly 130 | Gln | Asp | Thr | Val | Thr 135 | Val | Gly | Gly | Ile | Thr 140 | Val | Thr | Asn | Gln |
| Gln 145 | Phe | Gly | Leu | Ala | Thr 150 | Lys | Glu | Pro | Gly | Ser 155 | Phe | Phe | Ala | Thr | Ala 160 |
| Val | Phe | Asp | Gly | Ile 165 | Leu | Gly | Leu | Gly | Phe 170 | Pro | Ser | Ile | Glu | Ala 175 | Gly |
| Gly | Pro | Tyr | Thr 180 | Pro | Val | Phe | Asp | Asn 185 | Leu | Lys | Ser | Gln | Gly 190 | Leu | Ile |
| Asp | Ser | Pro 195 | Ala | Phe | Ser | Val | Tyr 200 | Leu | Asn | Ser | Asp | Ser 205 | Gly | Ala | Gly |
| Gly | Glu 210 | Ile | Ile | Phe | Gly | Gly 215 | Val | Asp | Pro | Ser | Lys 220 | Tyr | Thr | Gly | Ser |
| Leu 225 | Thr | Trp | Val | Pro | Val 230 | Thr | Ser | Gln | Gly | Tyr 235 | Trp | Gln | Ile | Thr | Leu 240 |
| Asp | Ser | Ile | Thr | Val 245 | Gly | Gly | Ser | Thr | Thr 250 | Phe | Cys | Ser | Ser | Gly 255 | Cys |
| Gln | Ala | Ile | Leu 260 | Asp | Thr | Gly | Thr | Ser 265 | Leu | Leu | Tyr | Gly | Pro 270 | Thr | Ser |
| Ile | Val | Ser 275 | Lys | Ile | Ala | Lys | Ala 280 | Val | Gly | Ala | Ser | Leu 285 | Ser | Glu | Tyr |
| Ser | Gly 290 | Glu | Tyr | Val | Ile | Asp 295 | Cys | Asp | Ser | Ile | Ser 300 | Ser | Leu | Pro | Asp |
| Ile 305 | Thr | Phe | Phe | Ile | Gly 310 | Gly | Ala | Lys | Ile | Thr 315 | Val | Pro | Pro | Ser | Ala 320 |
| Tyr | Val | Leu | Gln | Pro 325 | Ser | Ser | Gly | Gly | Ser 330 | Asp | Ile | Cys | Leu | Ser 335 | Gly |

Phe Gln Ser Asp Asp Ile Pro Gly Gly Pro Leu Trp Ile Leu Gly Asp 340 345 350

Val Phe Leu Arg Ser Ala Tyr Val Val Phe Asp Arg Asp Asn Asn Arg 355 360 365

Ile Gly Leu Ala Pro Ala 370

<210> 135

<211> 208

<212> PRT

<213> Mus musculus

<400> 135

Met Lys Val Thr Leu Val His Leu Leu Phe Met Met Leu Leu Leu 1 5 10 15

Leu Gly Leu Gly Leu Gly Leu Gly Leu His Met Ala Ala Ala 20 25 30

Val Leu Glu Asp Gln Pro Leu Asn Glu Phe Trp Pro Ser Asp Ser Gln
35 40 45

Asn Thr Glu Glu Gly Glu Gly Ile Trp Thr Thr Glu Gly Leu Ala Leu 50 55 60

Gly Tyr Lys Glu Met Ala Gln Pro Val Trp Pro Glu Glu Ala Val Leu
65 70 75 80

Ser Glu Asp Glu Val Gly Gly Ser Arg Met Leu Arg Ala Glu Pro Arg 85 90 95

Phe Gln Ser Lys Gln Asp Tyr Leu Lys Phe Asp Leu Ser Val Arg Asp
100 105 110

Cys Asn Thr Met Met Ala His Lys Ile Lys Glu Pro Asn Gln Ser Cys 115 120 125

Ile Asn Gln Tyr Thr Phe Ile His Glu Asp Pro Asn Thr Val Lys Ala 130 135 140

Val Cys Asn Gly Ser Leu Val Asp Cys Asp Leu Gln Gly Gly Lys Cys 145 150 155 160

Tyr Lys Ser Pro Arg Pro Phe Asp Leu Thr Leu Cys Lys Leu Ala Lys

165 170 175

Pro Gly Gln Val Thr Pro Asn Cys His Tyr Leu Thr Tyr Ile Thr Glu 180 185 190

Lys Ser Ile Phe Met Thr Cys Asn Asp Lys Arg Gln Leu Glu Thr Lys 195 200 205

<210> 136

<211> 149

<212> PRT

<213> Mus pahari

<400> 136

Met Gly Leu Glu Lys Ser Leu Ile Leu Phe Pro Leu Phe Val Leu Leu

1 5 10 15

Leu Gly Trp Val Gln Pro Ser Leu Gly Lys Glu Ser Ser Ala Gln Lys
20 25 30

Phe Glu Arg Gln His Met Asp Ser Ser Gly Ser Ser Asn Asn Ser Pro 35 40 45

Thr Tyr Cys Asn Gln Met Met Lys Ser Arg Ser Met Thr Lys Glu Ser 50 55 60

Cys Lys Pro Val Asn Thr Phe Val His Glu Pro Leu Glu Asp Val Gln 65 70 75 80

Ala Ile Cys Ser Gln Glu Asn Val Thr Cys Lys Asn Gly Asn Arg Asn 85 90 95

Cys Tyr Lys Ser Ser Ser Ala Leu His Ile Thr Asp Cys His Leu Lys
100 105 110

Gly Asn Ser Lys Tyr Pro Asn Cys Asn Tyr Asn Thr Asn Gln Tyr Gln
115 120 125

Lys His Ile Ile Val Ala Cys Asp Gly Asn Pro Tyr Val Pro Val His 130 135 140

Leu Asp Ala Thr Val 145 <211> 125 <212> PRT <213> Homo sapiens <400> 137 Xaa Lys Glu Ser Ala Ala Ala Lys Phe Glu Arg Gln His Met Asp Ser 10 15 Gly Asn Ser Pro Ser Ser Ser Ser Thr Tyr Cys Asn Gln Met Met Arg 20 25 Arg Arg Asn Met Thr Gln Gly Arg Cys Lys Pro Val Asn Thr Phe Val 40 His Glu Ser Leu Val Asp Val Gln Asn Val Cys Phe Gln Glu Lys Val 55 60 Thr Cys Lys Asn Gly Gln Gly Asn Cys Tyr Lys Ser Asn Ser Ser Met His Ile Thr Asp Cys Arg Leu Thr Asn Gly Ser Arg Tyr Pro Asn Cys 85 90 Ala Tyr Arg Thr Ser Pro Lys Glu Arg His Ile Ile Val Ala Cys Glu 100 105 Gly Ser Pro Tyr Val Pro Val His Phe Asp Ala Ser Val 115 120 125 <210> 138 <211> 128 <212> PRT <213> Presbytis entellus <400> 138 Gly Glu Ser Arg Ala Glu Lys Phe Gln Arg Gln His Met Asp Ser Gly 10 Ser Ser Pro Ser Ser Ser Thr Tyr Cys Asn Gln Met Met Lys Leu 25 Arg Asn Met Thr Gln Gly Ser Cys Lys Ser Val Asn Thr Phe Val His 35 40 45

<210> 137

Glu Pro Leu Val Asp Val Gln Asn Val Cys Phe Gln Glu Lys Val Thr

50 55 60

Cys Lys Asn Gly Gln Thr Asn Cys Phe Lys Ser Asn Ser Arg Met His
65 70 75 80

Ile Thr Glu Cys Arg Leu Thr Asn Gly Ser Lys Tyr Pro Asn Cys Ala 85 90 95

Tyr Gly Thr Ser Pro Lys Glu Arg His Ile Ile Val Ala Cys Glu Gly
100 105 110

Ser Pro Tyr Val Pro Val His Phe Asp Asp Ser Val Glu Asp Ser Thr 115 120 125

<210> 139

<211> 119

<212> PRT

<213> Iguana iguana

<400> 139

Gln Asp Trp Ser Ser Phe Gln Asn Lys His Ile Asp Tyr Pro Glu Thr
1 5 10 15

Ser Ala Ser Asn Pro Asn Ala Tyr Cys Asp Leu Met Met Gln Arg Arg 20 25 30

Asn Leu Asn Pro Thr Lys Cys Lys Thr Arg Asn Thr Phe Val His Ala $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Ser Pro Ser Glu Ile Gln Gln Val Cys Gly Ser Gly Gly Thr His Tyr 50 60

Glu Asp Asn Leu Tyr Asp Ser Asn Glu Ser Phe Asp Leu Thr Asp Cys
65 70 75 80

Lys Asn Val Gly Gly Thr Ala Pro Ser Ser Cys Lys Tyr Asn Gly Thr 85 90 95

Pro Gly Thr Lys Arg Ile Arg Ile Ala Cys Glu Asn Asn Gln Pro Val 100 105 110

His Phe Glu Leu Val Leu Ser

<210> 140 <211> 105 <212> PRT <213> Homo sapiens

<400> 140

His Val Asp Tyr Pro Gln Asn Asp Val Pro Val Pro Ala Arg Tyr Cys
1 5 10 15

Asn His Met Ile Ile Gln Arg Val Ile Arg Glu Pro Asp His Thr Cys
20 25 30

Lys Lys Glu His Val Phe Ile His Glu Arg Pro Arg Lys Ile Asn Gly 35 40 45

Ile Cys Ile Ser Pro Lys Lys Val Ala Cys Gln Asn Leu Ser Ala Ile 50 55 60

Phe Cys Phe Gln Ser Glu Thr Lys Phe Lys Met Thr Val Cys Gln Leu 65 70 75 80

Ile Glu Gly Thr Arg Tyr Pro Ala Cys Arg Tyr His Tyr Ser Pro Thr
85 90 95

Glu Gly Phe Val Leu Val Thr Cys Asp 100 105

<210> 141

<211> 99

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: RNAse_Pc,
 Pancreatic ribonuclease

<400> 141

His Ile Asp Ser Thr Pro Ser Ser Ala Ser Asp Asn Tyr Cys Asn Gln
1 5 10 15

Met Met Lys Arg Arg Asn Met Thr Gln Gly Arg Cys Lys Pro Val Asn 20 25 30

Thr Phe Val His Glu Ser Leu Ala Asp Val Lys Ala Val Cys Ser Gln 35 40 45

Lys Asn Val Thr Cys Lys Asn Gly Arg Thr Asn Cys His Gln Ser Asn 50 55 60

Ser Arg Phe Gln Leu Thr Asp Cys Arg Leu Thr Gly Gly Ser Lys Tyr 65 70 75 80

Pro Asn Cys Arg Tyr Lys Thr Thr Gln Ala Asn Lys His Ile Ile Val 85 90 95

Ala Cys Glu

<210> 142

<211> 93

<212> PRT

<213> Homo sapiens

<400> 142

Ala Arg Tyr Cys Asn His Met Ile Ile Gln Arg Val Ile Arg Glu Pro 1 5 10 15

Asp His Thr Cys Lys Lys Glu His Val Phe Ile His Glu Arg Pro Arg 20 25 30

Lys Ile Asn Gly Ile Cys Ile Ser Pro Lys Lys Val Ala Cys Gln Asn 35 40 45

Leu Ser Ala Ile Phe Cys Phe Gln Ser Glu Thr Lys Phe Lys Met Thr
50 55 60

Val Cys Gln Leu Ile Glu Gly Thr Arg Tyr Pro Ala Cys Arg Tyr His 65 70 75 80

Tyr Ser Pro Thr Glu Gly Phe Val Leu Val Thr Cys Asp 85 90

<210> 143

<211> 89

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: rnaseA, Pancreatic ribonuclease

<400> 143

Asp Asn Tyr Cys Asn Gln Met Met Lys Arg Arg Asn Met Thr Gln Gly

1 5 10 15

Arg Cys Lys Pro Val Asn Thr Phe Val His Glu Ser Leu Ala Asp Val 20 25 30

Lys Ala Val Cys Ser Gln Lys Asn Val Thr Cys Lys Asn Gly Gln Lys 35 40 45

Asn Cys Tyr Gln Ser Thr Ser Ser Phe Gln Leu Thr Asp Cys Arg Leu 50 55 60

Thr Gly Gly Ser Lys Tyr Pro Asn Cys Arg Tyr Arg Thr Thr Pro Gly 65 70 75 80

Asn Lys Arg Ile Ile Val Ala Cys Glu 85

<210> 144

<211> 698

<212> PRT

<213> Mus musculus

<400> 144

Met Glu Lys Tyr Glu Arg Ile Arg Val Val Gly Arg Gly Ala Phe Gly
1 5 10 15

Ile Val His Leu Cys Leu Arg Lys Ala Asp Gln Lys Leu Val Ile Leu 20 25 30

Lys Gln Ile Pro Val Glu Gln Met Thr Lys Glu Glu Arg Gln Ala Ala 35 40 45

Gln Asn Glu Cys Gln Val Leu Lys Leu Leu Asn His Pro Asn Val Ile
50 55 60

Glu Tyr Tyr Glu Asn Phe Leu Glu Asp Lys Ala Leu Met Ile Ala Met
65 70 75 80

Glu Tyr Ala Pro Gly Gly Thr Leu Ala Glu Phe Ile Gln Lys Arg Cys 85 90 95

Asn Ser Leu Leu Glu Glu Glu Thr Ile Leu His Phe Phe Val Gln Ile
100 105 110

Leu Leu Ala Leu His His Val His Thr His Leu Ile Leu His Arg Asp 115 120 125

| Leu | Lys 130 | Thr | Gln | Asn | Ile | Leu 135 | Leu | Asp | Lys | His | Arg 140 | Met | Val | Val | Lys |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile 145 | Gly | Asp | Phe | Gly | Ile 150 | Ser | Lys | Ile | Leu | Ser 155 | Ser | Lys | Ser | Lys | Ala 160 |
| Tyr | Thr | Val | Val | Gly 165 | Thr | Pro | Cys | Tyr | Ile 170 | Ser | Pro | Glu | Leu | Cys 175 | Glu |
| Gly | Lys | Pro | Tyr 180 | Asn | Gln | Lys | Ser | Asp 185 | Ile | Trp | Ala | Leu | Gly 190 | Cys | Val |
| Leu | Tyr | Glu 195 | Leu | Ala | Ser | Leu | Lys 200 | Arg | Ala | Phe | Glu | Ala 205 | Ala | Asn | Leu |
| Pro | Ala 210 | Leu | Val | Leu | Lys | Ile 215 | Met | Ser | Gly | Thr | Phe 220 | Ala | Pro | Ile | Ser |
| Asp 225 | Arg | Tyr | Ser | Pro | Glu 230 | Leu | Arg | Gln | Leu | Val 235 | Leu | Ser | Leu | Leu | Ser 240 |
| Leu | Glu | Pro | Ala | Gln 245 | Arg | Pro | Pro | Leu | Ser 250 | His | Ile | Met | Ala | Gln 255 | Pro |
| Leu | Cys | Ile | Arg 260 | Ala | Leu | Leu | Asn | Ile 265 | His | Thr | Asp | Val | Gly 270 | Ser | Val |
| Arg | Met | Arg 275 | Arg | Ala | Glu | Lys | Ser 280 | Leu | Thr | Pro | Gly | Pro 285 | Pro | Ile | Ala |
| Ser | Gly 290 | Ser | Thr | Gly | Ser | Arg 295 | Ala | Thr | Ser | Ala | Arg 300 | Cys | Arg | Gly | Val |
| Pro 305 | Arg | Gly | Pro | Val | Arg 310 | Pro | Ala | Ile | Pro | Pro 315 | Pro | Leu | Ser | Ser | Val 320 |
| Tyr | Ala | Trp | Gly | Gly 325 | Gly | Leu | Ser | Ser | Pro 330 | Leu | Arg | Leu | Pro | Met 335 | Leu |
| Asn | Thr | Glu | Val 340 | Val | Gln | Val | Ala | Ala 345 | Gly | Arg | Thr | Gln | Lys 350 | Ala | Gly |
| Val | Thr | Arg 355 | Ser | Gly | Arg | Leu | Ile 360 | Leu | Trp | Glu | Ala | Pro 365 | Pro | Leu | Gly |
| Ala | Gly 370 | Gly | Gly | Thr | Leu | Leu 375 | Pro | Gly | Ala | Val | Glu 380 | Leu | Pro | Gln | Pro |

| Gln 385 | Phe | Val | Ser | Arg | Phe 390 | Leu | Glu | Gly | Gln | Ser 395 | Gly | Val | Thr | Ile | Lys 400 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Val | Ala | Cys | Gly 405 | Asp | Leu | Phe | Thr | Ala 410 | Cys | Leu | Thr | Asp | Arg 415 | Gly |
| Ile | Ile | Met | Thr 420 | Phe | Gly | Ser | Gly | Ser 425 | Asn | Gly | Cys | Leu | Gly 430 | His | Gly |
| Asn | Leu | Thr 435 | Asp | Ile | Ser | Gln | Pro 440 | Thr | Ile | Val | Glu | Ala 445 | Leu | Leu | Gly |
| Tyr | Glu 450 | Met | Val | Gln | Val | Ala 455 | Cys | Gly | Ala | Ser | His 460 | Val | Leu | Ala | Leu |
| Ser 465 | Thr | Asp | Gly | Glu | Leu 470 | Phe | Ala | Trp | Gly | Arg 475 | Gly | Asp | Gly | Gly | Arg 480 |
| Leu | Gly | Leu | Gly | Thr 485 | Arg | Glu | Ser | His | Asn 490 | Cys | Pro | Gln | Gln | Val 495 | Pro |
| Val | Ala | Pro | Gly 500 | Gln | Glu | Ala | Gln | Arg 505 | Val | Val | Cys | Gly | Ile 510 | Asp | Ser |
| Ser | Met | Ile 515 | Leu | Thr | Ser | Pro | Gly 520 | Arg | Val | Leu | Ala | Cys 525 | Gly | Ser | Asn |
| Arg | Phe 530 | Asn | Lys | Leu | Gly | Leu 535 | Asp | His | Leu | Ser | Leu 540 | Asp | Glu | Glu | Pro |
| Val 545 | Pro | Tyr | Gln | Gln | Val 550 | Glu | Glu | Ala | Leu | Ser 555 | Phe | Thr | Pro | Leu | Gly 560 |
| Ser | Ala | Pro | Leu | Asp 565 | Gln | Glu | Pro | Leu | Leu 570 | Cys | Val | Asp | Leu | Gly 575 | Thr |
| Ala | His | Ser | Ala 580 | Ala | Ile | Thr | Ala | Ser 585 | Gly | Asp | Cys | Tyr | Thr 590 | Phe | Gly |
| Ser | Asn | Gln 595 | His | Gly | Gln | Leu | Gly 600 | Thr | Ser | Ser | Arg | Arg 605 | Val | Ser | Arg |
| Ala | Pro 610 | Cys | Arg | Val | Gln | Gly 615 | Leu | Glu | Gly | Ile | Lys 620 | Met | Val | Met | Val |
| Ala 625 | Cys | Gly | Asp | Ala | Phe 630 | Thr | Val | Ala | Val | Gly 635 | Ala | Glu | Gly | Glu | Val 640 |

Tyr Ser Trp Gly Lys Gly Thr Arg Gly Arg Leu Gly Arg Arg Asp Glu 645 650 655

Asp Ala Gly Leu Pro Arg Pro Val Gln Leu Asp Glu Thr His Pro Tyr 660 665 670

Met Val Thr Ser Val Ser Cys Cys His Gly Asn Thr Leu Leu Ala Val 675 680 685

Arg Ser Val Thr Asp Glu Pro Val Pro Pro 690 695

<210> 145

<211> 291

<212> PRT

<213> Mus musculus

<400> 145

Met Glu Lys Tyr Glu Arg Ile Arg Val Val Gly Arg Gly Ala Leu Gly

1 5 10 15

Ile Val His Leu Cys Leu Arg Lys Ala Asp Gln Lys Leu Val Ile Leu 20 25 30

Lys Gln Ile Pro Val Glu Gln Met Thr Lys Glu Glu Arg Gln Ala Ala 35 40 45

Gln Asn Glu Cys Gln Val Leu Lys Leu Leu Asn His Pro Asn Val Ile 50 55 60

Glu Tyr Tyr Glu Asn Phe Leu Glu Asp Lys Ala Leu Met Ile Ala Met
65 70 75 80

Glu Tyr Ala Pro Gly Gly Thr Leu Ala Glu Phe Ile Gln Lys Arg Cys
85 90 95

Asn Ser Leu Leu Glu Glu Glu Thr Ile Leu His Phe Phe Val Gln Ile 100 105 110

Leu Leu Ala Leu His His Val His Thr His Leu Ile Leu His Arg Asp 115 120 125

Leu Lys Thr Gln Asn Ile Leu Leu Asp Lys His Arg Met Val Val Lys
130 135 140

Ile Gly Asp Phe Gly Ile Ser Lys Ile Leu Ser Ser Lys Ser Lys Ala

Tyr Thr Val Val Gly Thr Pro Cys Tyr Ile Ser Pro Glu Leu Cys Glu 165 170 175

Gly Lys Pro Tyr Asn Gln Lys Ser Asp Ile Trp Ala Leu Gly Cys Val 180 185 190

Leu Tyr Glu Leu Ala Ser Leu Lys Arg Ala Phe Glu Ala Ala Asn Leu 195 200 205

Pro Ala Leu Val Leu Lys Ile Met Ser Gly Thr Phe Ala Pro Ile Ser 210 215 220

Asp Arg Tyr Ser Pro Glu Leu Arg Gln Leu Val Leu Ser Leu Leu Ser 225 230 235 240

Leu Glu Pro Ala Gln Gly Pro Pro Leu Ser His Ile Met Ala Gln Pro 245 250 255

Leu Cys Ile Arg Ala Leu Leu Asn Ile His Thr Asp Val Gly Ser Val 260 265 270

Arg Met Arg Arg Pro Val Gln Gly Asp Gly Ser Trp Gly Gly His Pro 275 280 285

Val Arg Thr 290

<210> 146

<211> 696

<212> PRT

<213> Danio rerio

<400> 146

Met Glu Lys Tyr Glu Lys Thr Lys Val Val Gly Arg Gly Ala Phe Gly
1 5 10 15

Ile Val His Leu Cys Arg Arg Arg Thr Asp Ser Ala Leu Val Ile Leu 20 25 30

Lys Glu Ile Pro Val Glu Gln Met Thr Arg Asp Glu Arg Leu Ala Ala 35 40 45

Gln Asn Glu Cys Gln Val Leu Lys Leu Leu Ser His Pro Asn Ile Ile 50 55 60

| Glu T 65 | ſyr | Tyr | Glu | Asn | Phe 70 | Leu | Glu | Asp | Lys | Ala 75 | Leu | Met | Ile | Ala | Met 80 |
|--------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu T | ſyr | Ala | Pro | Gly 85 | Gly | Thr | Leu | Ala | Asp 90 | Tyr | Ile | Gln | Lys | Arg 95 | Cys |
| Asn S | Ser | Leu | Leu 100 | Asp | Glu | Asp | Thr | Ile 105 | Leu | His | Ser | Phe | Val 110 | Gln | Ile |
| Leu I | Leu | Ala 115 | Leu | Tyr | His | Val | His 120 | Asn | Lys | Leu | Ile | Leu 125 | His | Arg | Asp |
| Leu I | r 7 | Thr | Gln | Asn | Ile | Leu 135 | Leu | Asp | Lys | His | Gln 140 | Met | Ile | Val | Lys |
| Ile G 145 | Sly | Asp | Phe | Gly | Ile 150 | Ser | Lys | Ile | Leu | Val 155 | Ser | Lys | Ser | Lys | Ala 160 |
| Tyr T | ľhr | Val | Val | Gly 165 | Thr | Pro | Cys | Tyr | Ile 170 | Ser | Pro | Glu | Leu | Cys 175 | Glu |
| Gly I | ùуs | Pro | Tyr 180 | Asn | Gln | Lys | Ser | Asp 185 | Ile | Trp | Ala | Leu | Gly 190 | Cys | Val |
| Leu T | ſyr | Glu 195 | Leu | Ala | Ser | Leu | Lys 200 | Arg | Ala | Phe | Glu | Ala 205 | Ala | Asn | Leu |
| Pro A | Ala 210 | Leu | Val | Leu | Lys | Ile 215 | Met | Ser | Gly | Thr | Phe 220 | Ala | Pro | Ile | Ser |
| Asp A 225 | Arg | Tyr | Ser | Pro | Glu 230 | Leu | Arg | Gln | Leu | Ile 235 | Leu | Asn | Met | Leu | Asn 240 |
| Leu A | Asp | Pro | Ser | Lys 245 | Arg | Pro | Gln | Leu | Asn 250 | Glu | Ile | Met | Ala | His 255 | Ala |
| Ile C | Cys | Ile | Arg 260 | Pro | Leu | Leu | Asn | Leu 265 | Tyr | Thr | Asp | Ile | Gly 270 | Asn | Val |
| Lys M | 1et | Arg 275 | Arg | Ile | Glu | Lys | Pro 280 | Leu | Ser | Asn | Val | Gln 285 | Ala | Gly | Pro |
| His G | 31y 290 | Arg | Pro | Gly | Gly | Trp 295 | Ile | Thr | Ser | Thr | Arg 300 | Thr | Arg | Gly | Gly |
| Leu S 305 | Ser | Ser | Leu | Thr | Ser 310 | Ser | Lys | Met | Met | His 315 | Pro | Leu | Pro | Leu | Phe 320 |

Ser Val Tyr Thr Trp Gly Ser Gly Ile Ser Thr Pro Leu Arg Leu Pro Met Leu Asn Thr Glu Val Ile Gln Val Ser Leu Gly Arg Thr Gln Lys Met Gly Val Thr Lys Ser Arg Leu Ile Thr Trp Glu Ala Pro Ser Val Gly Ser Gly Glu Pro Thr Leu Pro Gly Ala Val Glu Gln Met Gln Pro Gln Phe Ile Ser Arg Phe Leu Glu Gly Gln Ser Gly Val Thr Ile Lys Ser Val Ser Cys Gly Asp Leu Phe Thr Thr Cys Leu Thr Asp Arg Gly Ile Ile Met Thr Phe Gly Ser Gly Ser Asn Gly Cys Leu Gly His Gly Asn Phe Asn Asp Val Thr Gln Pro Lys Ile Val Glu Ala Leu Leu Gly Tyr Glu Leu Val Gln Val Ser Cys Gly Ala Ser His Val Leu Ala Val Thr Asn Glu Arg Glu Val Phe Ser Trp Gly Arg Gly Asp Asn Gly Arg Leu Gly Leu Ala Thr Gln Asp Ser His Asn Cys Pro Gln Gln Val Ser Leu Pro Ala Asp Phe Glu Ala Gln Arg Val Leu Cys Gly Val Asp Cys Ser Met Ile Met Ser Thr Gln His Gln Ile Leu Ala Cys Gly Asn Asn Arg Phe Asn Lys Leu Gly Leu Asp Lys Val Ser Gly Thr Glu Glu Pro Ser Ser Phe Cys Gln Val Glu Glu Val His Leu Phe Gln Leu Val Gln Ser Ala Pro Leu Asn Thr Glu Lys Ile Val Tyr Ile Asp Ile Gly Thr

Ala His Ser Val Ala Val Thr Glu Lys Gly Gln Cys Phe Thr Phe Gly Ser Asn Gln His Gly Gln Leu Gly Cys Ser His Arg Arg Ser Ser Arg Val Pro Tyr Gln Val Ser Gly Leu Gln Gly Ile Thr Met Ala Ala Cys Gly Asp Ala Phe Thr Leu Ala Ile Gly Ala Glu Gly Glu Val Tyr Thr Trp Gly Lys Gly Ala Arg Gly Arg Leu Gly Arg Lys Glu Glu Asp Phe Gly Ile Pro Lys Pro Val Gln Leu Asp Glu Ser His Ala Phe Thr Val Thr Ser Val Ala Cys Cys His Gly Asn Thr Leu Leu Ala Val Lys Pro Phe Phe Glu Glu Pro Gly Pro Lys <210> 147 <211> 357 <212> PRT <213> Caenorhabditis elegans <400> 147 Met Asp Asn Tyr Glu Lys Val Arg Val Gly Arg Gly Ala Phe Gly Val Cys Trp Leu Cys Arg Gly Lys Asn Asp Ala Ser His Gln Lys Val Ile Ile Lys Leu Ile Asn Thr His Gly Met Thr Glu Lys Glu Glu Asn Ser Ile Gln Ser Glu Val Asn Leu Leu Lys Lys Val Gln His Pro Leu Ile Ile Gly Tyr Ile Asp Ser Phe Ile Met Asp Asn Gln Leu Gly Ile Val Met Gln Tyr Ala Glu Gly Gly Thr Leu Glu Arg Leu Ile Asn Asp

Gln Arg Ala Ile Lys Asp Ser Asn Met Arg Glu Tyr Phe Pro Glu Lys Thr Val Leu Asp Tyr Phe Thr Gln Ile Leu Ile Ala Leu Asn His Met His Gln Lys Asn Ile Val His Arg Asp Leu Lys Pro Gln Asn Ile Leu Met Asn Arg Arg Lys Thr Val Leu Lys Leu Ser Asp Phe Gly Ile Ser Lys Glu Leu Gly Thr Lys Ser Ala Ala Ser Thr Val Ile Gly Thr Pro Asn Tyr Leu Ser Pro Glu Ile Cys Glu Ser Arg Pro Tyr Asn Gln Lys Ser Asp Met Trp Ser Leu Gly Cys Val Leu Tyr Glu Leu Leu Gln Leu Glu Arg Ala Phe Asp Gly Glu Asn Leu Pro Ala Ile Val Met Lys Ile Thr Arg Ser Lys Gln Asn Pro Leu Gly Asp His Val Ser Asn Asp Val Lys Met Leu Val Glu Asn Leu Leu Lys Thr His Thr Asp Lys Arg Pro Asp Val Ser Gln Leu Leu Ser Asp Pro Leu Val Leu Pro Tyr Leu Ile Ser Ile His Cys Asp Leu Gly Arg Ile Glu Pro Pro Pro Thr Asp Lys Arg Lys Pro Ser Ala Ser Leu Ser Ser Arg Leu Arg Thr Tyr Pro Thr Gln Ser Thr Leu Arg Pro Tyr Ser Leu Ser Ser Asn Ala Pro Thr Thr His Leu Thr Gln Leu Thr Pro Met Pro Ser His Ile Asp Ser Gly Phe Phe Ser Ser Gly Arg Thr Ser Asn Gln Arg Thr Gln Ser Arg Ser Gln

Val His Ser Lys Tyr 355

<210> 148

<211> 841

<212> PRT

<213> Drosophila melanogaster

<400> 148

Met Lys Lys Phe Arg Ala Lys Ala Ser Ser Leu Pro Ile Phe Asn Gly
1 5 10 15

Arg Ile Thr Asp Ala Thr Thr Leu Thr Thr Ser Ser Leu Gln Leu Pro 20 25 30

Leu Gly Gln Asn Thr Gln Arg Lys Gln Ser Thr Cys Thr Arg Val Leu 35 40 45

Pro Thr Val Phe Thr Ile Thr Asp Gly Thr Thr Gly Ala Ala Ser Thr 50 55 60

Ser Leu Ala Glu Ala Met Ser Ser Ser Lys Ala Gln Met Pro Asn Arg
65 70 75 80

Gln Glu Ser Leu Leu Gln Leu Ser Val Pro Arg Glu Thr Gly Val Gly
85 90 95

Val Ala Gly Pro Glu Leu Ala Asn Tyr Glu Lys Val Arg Val Val Gly
100 105 110

Gln Gly Ser Phe Gly Ile Ala Ile Leu Tyr Arg Arg Lys Ser Asp Gly
115 120 125

His Gln Ile Val Phe Lys Gln Ile Asn Leu Ser Glu Leu Ser Pro Pro 130 135 140

Gly Arg Asp Leu Ala Met Asn Glu Val Asp Val Phe Ser Lys Leu His 145 150 155 160

His Pro Asn Ile Val Ser Tyr Leu Gly Ser Phe Ile Lys Asp Asn Thr 165 170 175

Leu Leu Ile Glu Met Glu Tyr Ala Asp Gly Gly Thr Leu Ala Gln Ile 180 185 190

Ile Ala Glu Arg Gln Gly Lys Leu His Phe Pro Glu Arg Tyr Ile Ile

| 195 | 200 | 205 |
|-----|-----|-----|
| | | |

Ala Val Phe Glu Gln Ile Ser Ser Ala Ile Asn Tyr Met His Ser Glu Asn Ile Leu His Arg Asp Leu Lys Thr Ala Asn Val Phe Leu Asn Arg Arg Gly Ile Val Lys Ile Gly Asp Phe Gly Ile Ser Lys Ile Met Asn Thr Lys Ile His Ala Gln Thr Val Leu Gly Thr Pro Tyr Tyr Phe Ser Pro Glu Met Cys Glu Gly Lys Glu Tyr Asp Asn Lys Ser Asp Ile Trp Ala Leu Gly Cys Ile Leu Gly Glu Met Cys Cys Leu Lys Lys Thr Phe Ala Ala Ser Asn Leu Ser Glu Leu Val Thr Lys Ile Met Ala Gly Asn Tyr Thr Pro Val Pro Ser Gly Tyr Thr Ser Gly Leu Arg Ser Leu Met Ser Asn Leu Leu Gln Val Glu Ala Pro Arg Arg Pro Thr Ala Ser Glu Val Leu Val Tyr Trp Ile Pro Leu Ile Phe Arg Ser Leu Gly Lys Asn Lys Gly Tyr Ser Tyr Glu Asp Asp Val Gly Gly Pro Gly Ser Asp Gln Leu Thr Ala Pro Val Pro Ala Ala Ala Tyr Ser Asn Val Ser Met Glu Leu Glu Leu Pro Thr Ala Gln Thr Glu Thr Lys Gln Leu Met Ile Ala Asp Thr Ala Ala Pro His Glu Ile Leu Glu Lys Arg Ser Val Leu Tyr Gln Leu Lys Ala Phe Gly Thr Cys Phe Ser Met Ala Pro Ile Gln Leu

Pro Pro Lys Ala Val Ile Val Asp Val Ala Met Ser Asp Ser His Phe

Val Val Val Asn Glu Asp Gly Ser Ala Tyr Ala Trp Gly Glu Gly Thr His Gly Gln Leu Gly Leu Thr Ala Leu Glu Ala Trp Lys His Tyr Pro Ser Arg Met Glu Ser Val Arg Asn Tyr His Val Val Ser Ala Cys Ala Gly Asp Gly Phe Thr Ile Leu Val Thr Gln Ala Gly Ser Leu Leu Ser Cys Gly Ser Asn Ala His Leu Ala Leu Gly Gln Asp Glu Gln Arg Asn Tyr His Ser Pro Lys Leu Ile Ala Arg Leu Ala Asp Val Arg Val Glu Gln Val Ala Ala Gly Leu Gln His Val Leu Ala Leu Ser Arg Glu Gly Ala Val Tyr Val Trp Gly Thr Ser Thr Cys Gly Ala Leu Gly Leu Gly Asn Tyr Gln Gln Gln Lys Phe Pro Gln Lys Ile Leu Leu Ser His Val Lys Thr Lys Pro Ser Lys Ile Tyr Cys Gly Pro Asp Thr Ser Ala Val Leu Phe Ala Asn Gly Glu Leu His Val Cys Gly Ser Asn Asp Tyr Asn Lys Leu Gly Phe Gln Arg Ser Ala Lys Ile Thr Ala Phe Lys Lys Val Gln Leu Pro His Lys Val Thr Gln Ala Cys Phe Ser Ser Thr His Ser Val Phe Leu Val Glu Gly Gly Tyr Val Tyr Thr Met Gly Arg Asn Ala Glu Gly Gln Arg Gly Ile Arg His Cys Asn Ser Val Asp His Pro

Thr Leu Val Asp Ser Val Lys Ser Arg Tyr Ile Val Lys Ala Asn Cys

| 705 | 710 | 715 | 720 |
|-----|-----|-----|-----|
| /03 | 710 | 113 | 120 |

Ser Asp Gln Cys Thr Ile Val Ala Ser Glu Asp Asn Ile Ile Thr Val 725 730 735

Trp Gly Thr Arg Asn Gly Leu Pro Gly Ile Gly Ser Thr Asn Cys Gly
740 745 750

Leu Gly Leu Gln Ile Cys Thr Pro Asn Met Glu Leu Glu Leu Gly Asn 755 760 765

Asn Thr Ala Ala Phe Thr Asn Phe Leu Ala Ser Val Tyr Lys Ser Glu 770 775 780

Leu Ile Leu Glu Pro Val Asp Ile Leu Ala Leu Phe Ser Ser Lys Glu 785 790 795 800

Gln Cys Asp Arg Gly Tyr Tyr Val Gln Val His Asp Val Tyr Pro Leu 805 810 815

Ala His Ser Val Leu Val Leu Val Asp Thr Thr Thr Pro Leu Ile Ser 820 825 830

Ser Tyr Glu Gly Asp Tyr Pro His Leu 835 840

<210> 149

<211> 253

<212> PRT

<213> Homo sapiens

<400> 149

Tyr Glu Arg Ile Arg Val Val Gly Arg Gly Ala Phe Gly Ile Val His

1 5 10 15

Leu Cys Leu Arg Lys Ala Asp Gln Lys Leu Val Ile Ile Lys Gln Ile 20 25 30

Pro Val Glu Gln Met Thr Lys Glu Glu Arg Gln Ala Ala Gln Asn Glu 35 40 45

Cys Gln Val Leu Lys Leu Leu Asn His Pro Asn Val Ile Glu Tyr Tyr 50 55 60

Glu Asn Phe Leu Glu Asp Lys Ala Leu Met Thr Ala Met Glu Tyr Ala 65 70 75 80

Pro Gly Gly Thr Leu Ala Glu Phe Ile Gln Lys Arg Cys Asn Ser Leu Leu Glu Glu Glu Thr Ile Leu His Phe Phe Val Gln Ile Leu Leu Ala Leu His His Val His Thr His Leu Ile Leu His Arg Asp Leu Lys Thr Gln Asn Ile Leu Leu Asp Lys His Arg Met Val Val Lys Ile Gly Asp Phe Gly Ile Ser Lys Ile Leu Ser Ser Lys Ser Lys Ala Tyr Thr Val Val Gly Thr Pro Cys Tyr Ile Ser Pro Glu Leu Cys Glu Gly Lys Pro Tyr Asn Gln Lys Ser Asp Ile Trp Ala Leu Gly Cys Val Leu Tyr Glu Leu Ala Ser Leu Lys Arg Ala Phe Glu Ala Ala Asn Leu Pro Ala Leu Val Leu Lys Ile Met Ser Gly Thr Phe Ala Pro Ile Ser Asp Arg Tyr Ser Pro Glu Leu Arg Gln Leu Val Leu Ser Leu Ser Leu Glu Pro Ala Gln Arg Pro Pro Leu Ser His Ile Met Ala Gln Pro <210> 150 <211> 254 <212> PRT <213> Artificial Sequence

<220>

<400> 150

Tyr Glu Leu Leu Glu Val Leu Gly Lys Gly Ala Phe Gly Lys Val Tyr

1 5 10 15

Leu Ala Arg Asp Lys Lys Thr Gly Lys Leu Val Ala Ile Lys Val Ile

- Lys Lys Glu Lys Leu Lys Lys Lys Arg Glu Arg Ile Leu Arg Glu
 35 40 45
- Ile Lys Ile Leu Lys Lys Leu Asp His Pro Asn Ile Val Lys Leu Tyr 50 55 60
- Asp Val Phe Glu Asp Asp Asp Lys Leu Tyr Leu Val Met Glu Tyr Cys
 65 70 75 80
- Glu Gly Gly Asp Leu Phe Asp Leu Leu Lys Lys Arg Gly Arg Leu Ser 85 90 95
- Glu Asp Glu Ala Arg Phe Tyr Ala Arg Gln Ile Leu Ser Ala Leu Glu 100 105 110
- Tyr Leu His Ser Gln Gly Ile Ile His Arg Asp Leu Lys Pro Glu Asn 115 120 125
- Ile Leu Leu Asp Ser Asp Gly His Val Lys Leu Ala Asp Phe Gly Leu 130 135 140
- Ala Lys Gln Leu Asp Ser Gly Gly Thr Leu Leu Thr Thr Phe Val Gly
 145 150 155 160
- Thr Pro Glu Tyr Met Ala Pro Glu Val Leu Leu Gly Lys Gly Tyr Gly
 165 170 175
- Lys Ala Val Asp Ile Trp Ser Leu Gly Val Ile Leu Tyr Glu Leu Leu 180 185 190
- Thr Gly Lys Pro Pro Phe Pro Gly Asp Asp Gln Leu Leu Ala Leu Phe 195 200 205
- Lys Lys Ile Gly Lys Pro Pro Pro Pro Pro Pro Pro Pro Glu Trp Lys 210 215 220
- Ile Ser Pro Glu Ala Lys Asp Leu Ile Lys Lys Leu Leu Val Lys Asp 225 230 235 240
- Pro Glu Lys Arg Leu Thr Ala Glu Glu Ala Leu Glu His Pro 245 250

<210> 151

<211> 254

<212> PRT

| <213 | 3> A1 | ctifi | icial | l Sed | queno | ce | | | | | | | | | |
|--------------|--------------|------------|----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| <220 <223 | 3> De | | iptio in ki | | | | | _ | | e: p} | cinas | se, | | | |
| |)> 15 Glu | | Gly | Glu 5 | Lys | Leu | Gly | Ser | Gly 10 | Ala | Phe | Gly | Lys | Val 15 | Tyr |
| Lys | Gly | Lys | His 20 | Lys | Asp | Thr | Gly | Glu 25 | Ile | Val | Ala | Ile | Lys 30 | Ile | Leu |
| Lys | Lys | Arg 35 | Ser | Leu | Ser | Glu | Lys 40 | Lys | Lys | Arg | Phe | Leu 45 | Arg | Glu | Ile |
| Gln | Ile 50 | Leu | Arg | Arg | Leu | Ser 55 | His | Pro | Asn | Ile | Val 60 | Arg | Leu | Leu | Gly |
| Val 65 | Phe | Glu | Glu | Asp | Asp 70 | His | Leu | Tyr | Leu | Val 75 | Met | Glu | Tyr | Met | Glu 80 |
| Gly | Gly | Asp | Leu | Phe 85 | Asp | Tyr | Leu | Arg | Arg 90 | Asn | Gly | Leu | Leu | Leu 95 | Ser |
| Glu | Lys | Glu | Ala 100 | Lys | Lys | Ile | Ala | Leu 105 | Gln | Ile | Leu | Arg | Gly 110 | Leu | Glu |
| Tyr | Leu | His 115 | Ser | Arg | Gly | Ile | Val 120 | His | Arg | Asp | Leu | Lys 125 | Pro | Glu | Asn |
| Ile | Leu 130 | Leu | Asp | Glu | Asn | Gly 135 | Thr | Val | Lys | Ile | Ala 140 | Asp | Phe | Gly | Leu |
| Ala 145 | Arg | Lys | Leu | Glu | Ser 150 | Ser | Ser | Tyr | Glu | Lys 155 | Leu | Thr | Thr | Phe | Val 160 |
| Gly | Thr | Pro | Glu | Tyr 165 | Met | Ala | Pro | Glu | Val 170 | Leu | Glu | Gly | Arg | Gly 175 | Tyr |
| Ser | Ser | Lys | Val 180 | Asp | Val | Trp | Ser | Leu 185 | Gly | Val | Ile | Leu | Tyr 190 | Glu | Leu |
| Leu | Thr | Gly 195 | Lys | Leu | Pro | Phe | Pro 200 | Gly | Ile | Asp | Pro | Leu 205 | Glu | Glu | Leu |

Phe Arg Ile Lys Glu Arg Pro Arg Leu Arg Leu Pro Leu Pro Pro Asn

Cys Ser Glu Glu Leu Lys Asp Leu Ile Lys Lys Cys Leu Asn Lys Asp 225 230 235 240

Pro Glu Lys Arg Pro Thr Ala Lys Glu Ile Leu Asn His Pro 245 250

<210> 152

<211> 245

<212> PRT

<213> Homo sapiens

<400> 152

Arg Val Val Gly Arg Gly Ala Phe Gly Ile Val His Leu Cys Leu Arg
1 5 10 15

Lys Ala Asp Gln Lys Leu Val Ile Ile Lys Gln Ile Pro Val Glu Gln 20 25 30

Met Thr Lys Glu Glu Arg Gln Ala Ala Gln Asn Glu Cys Gln Val Leu 35 40 45

Lys Leu Leu Asn His Pro Asn Val Ile Glu Tyr Tyr Glu Asn Phe Leu 50 55 60

Glu Asp Lys Ala Leu Met Thr Ala Met Glu Tyr Ala Pro Gly Gly Thr
65 70 75 80

Leu Ala Glu Phe Ile Gln Lys Arg Cys Asn Ser Leu Leu Glu Glu Glu 85 90 95

Thr Ile Leu His Phe Phe Val Gln Ile Leu Leu Ala Leu His His Val
100 105 110

His Thr His Leu Ile Leu His Arg Asp Leu Lys Thr Gln Asn Ile Leu 115 120 125

Leu Asp Lys His Arg Met Val Val Lys Ile Gly Asp Phe Gly Ile Ser 130 135 140

Lys Ile Leu Ser Ser Lys Ser Lys Ala Tyr Thr Val Val Gly Thr Pro 145 150 155 160

Cys Tyr Ile Ser Pro Glu Leu Cys Glu Gly Lys Pro Tyr Asn Gln Lys 165 170 175

Ser Asp Ile Trp Ala Leu Gly Cys Val Leu Tyr Glu Leu Ala Ser Leu

13.11

180 185 190

Lys Arg Ala Phe Glu Ala Ala Asn Leu Pro Ala Leu Val Leu Lys Ile 195 200 205

Met Ser Gly Thr Phe Ala Pro Ile Ser Asp Arg Tyr Ser Pro Glu Leu 210 215 220

Arg Gln Leu Val Leu Ser Leu Leu Ser Leu Glu Pro Ala Gln Arg Pro 225 230 235 240

Pro Leu Ser His Ile 245

<210> 153

<211> 250

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: TyrKc, Tyrosine kinase domain

<400> 153

Lys Lys Leu Gly Glu Gly Ala Phe Gly Glu Val Tyr Lys Gly Thr Leu
1 5 10 15

Lys Gly Lys Gly Gly Val Glu Val Glu Val Ala Val Lys Thr Leu Lys
20 25 30

Glu Asp Ala Ser Glu Gln Gln Ile Glu Glu Phe Leu Arg Glu Ala Arg 35 40 45

Leu Met Arg Lys Leu Asp His Pro Asn Ile Val Lys Leu Gly Val 50 55 60

Cys Thr Glu Glu Glu Pro Leu Met Ile Val Met Glu Tyr Met Glu Gly 65 70 75 80

Gly Asp Leu Leu Asp Tyr Leu Arg Lys Asn Arg Pro Lys Glu Leu Ser 85 90 95

Leu Ser Asp Leu Leu Ser Phe Ala Leu Gln Ile Ala Arg Gly Met Glu
100 105 110

Tyr Leu Glu Ser Lys Asn Phe Val His Arg Asp Leu Ala Ala Arg Asn 115 120 125 Cys Leu Val Gly Glu Asn Lys Thr Val Lys Ile Ala Asp Phe Gly Leu 130 135 Ala Arg Asp Leu Tyr Asp Asp Tyr Tyr Arg Lys Lys Ser Pro 150 155 145 160 Arg Leu Pro Ile Arg Trp Met Ala Pro Glu Ser Leu Lys Asp Gly Lys 165 170 Phe Thr Ser Lys Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu 180 185 Ile Phe Thr Leu Gly Glu Ser Pro Tyr Pro Gly Met Ser Asn Glu Glu 200 205 Val Leu Glu Tyr Leu Lys Lys Gly Tyr Arg Leu Pro Gln Pro Pro Asn 215 Cys Pro Asp Glu Ile Tyr Asp Leu Met Leu Gln Cys Trp Ala Glu Asp 230 235 Pro Glu Asp Arg Pro Thr Phe Ser Glu Leu 245 <210> 154 <211> 488 <212> PRT <213> Mus musculus <400> 154 Met Arg Ser Gly Ala Glu Arg Arg Gly Ser Ser Ala Ala Ala Pro Pro 1 10 Ser Ser Pro Pro Pro Gly Arg Ala Arg Pro Ala Gly Ser Glu Val Ser 20 25 Pro Ala Leu Pro Pro Pro Ala Ala Ser Gln Pro Arg Ala Arg Asp Ala Gly Asp Ala Arg Ala Gln Pro Arg Pro Leu Phe Gln Trp Ser Lys Trp 55 60 Lys Lys Arg Met Ser Met Ser Ser Ile Ser Ser Gly Ser Ala Arg Arg 75

Pro Val Phe Asp Asp Lys Glu Asp Val Asn Phe Asp His Phe Gln Ile

| Leu | Arg | Ala | Ile 100 | Gly | Lys | Gly | Ser | Phe 105 | Gly | Lys | Val | Cys | Ile 110 | Val | Glr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Lys | Arg | Asp 115 | Thr | Glu | Lys | Met | Tyr 120 | Ala | Met | Lys | Tyr | Met 125 | Asn | Lys | Glr |
| Gln | Cys 130 | Ile | Glu | Arg | Asp | Glu 135 | Val | Arg | Asn | Val | Phe 140 | Arg | Glu | Leu | Glu |
| Ile 145 | Leu | Gln | Glu | Ile | Glu 150 | His | Val | Phe | Leu | Val 155 | Asn | Leu | Trp | Tyr | Ser 160 |
| Phe | Gln | Asp | Glu | Glu 165 | Asp | Met | Phe | Met | Val 170 | Val | Asp | Leu | Leu | Leu 175 | Gly |
| Gly | Asp | Leu | Arg 180 | Tyr | His | Leu | Gln | Gln 185 | Asn | Val | Gln | Phe | Ser 190 | Glu | Asp |
| Thr | Val | Arg 195 | Leu | Tyr | Ile | Cys | Glu 200 | Met | Ala | Leu | Ala | Leu 205 | Asp | Tyr | Leı |
| Arg | Ser 210 | Gln | His | Ile | Ile | His 215 | Arg | Asp | Val | Lys | Pro 220 | Asp | Asn | Ile | Leı |
| Leu 225 | Asp | Glu | Gln | Gly | His 230 | Ala | His | Leu | Thr | Asp 235 | Phe | Asn | Ile | Ala | Th: |
| Ile | Ile | Lys | Asp | Gly 245 | Glu | Arg | Ala | Thr | Ala 250 | Leu | Ala | Gly | Thr | Lys 255 | Pro |
| Tyr | Met | Ala | Pro 260 | Glu | Ile | Phe | His | Ser 265 | Phe | Val | Asn | Gly | Gly 270 | Thr | Gl |
| Tyr | Ser | Phe 275 | Glu | Val | Asp | Trp | Trp 280 | Ser | Val | Gly | Val | Met 285 | Ala | Tyr | Glu |
| Leu | Leu 290 | Arg | Gly | Trp | Arg | Pro 295 | Tyr | Asp | Ile | His | Ser 300 | Ser | Asn | Ala | Va] |
| Glu 305 | Ser | Leu | Val | Gln | Leu 310 | Phe | Ser | Thr | Val | Ser 315 | Val | Gln | Tyr | Val | Pro 320 |
| Thr | Trp | Ser | Lys | Glu 325 | Met | Val | Ala | Leu | Leu 330 | Arg | Lys | Leu | Leu | Thr 335 | Va] |

Asn Pro Glu His Arg Phe Ser Ser Leu Gln Asp Met Gln Thr Ala Pro

| 340 | 345 | 350 |
|-----|-----|-----|
| | | |

| Ser | Leu | Ala | His | Val | Leu | Trp | Asp | Asp | Leu | Ser | Glu | Lys | Lys | Val | Glu |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | 355 | | | | | 360 | | | | | 365 | | | |

Pro Gly Phe Val Pro Asn Lys Gly Arg Leu His Cys Asp Pro Thr Phe 370 380

Glu Leu Glu Glu Met Ile Leu Glu Ser Arg Pro Leu His Lys Lys 385 390 395 400

Lys Arg Leu Ala Lys Asn Lys Ser Arg Asp Ser Ser Arg Asp Ser Ser 405 410 415

Gln Ser Glu Asn Asp Tyr Leu Gln Asp Cys Leu Asp Ala Ile Gln Gln 420 425 430

Asp Phe Val Ile Phe Asn Arg Glu Lys Leu Lys Arg Ser Gln Glu Leu 435 440 445

Met Ser Glu Pro Pro Pro Gly Pro Glu Thr Ser Asp Met Thr Asp Ser 450 455 460

Thr Ala Asp Ser Glu Ala Glu Pro Thr Ala Leu Pro Met Cys Gly Ser 465 470 475 480

Ile Cys Pro Ser Ser Gly Ser Ser 485

<210> 155

<211> 369

<212> PRT

<213> Homo sapiens

<400> 155

Met Tyr Ala Met Lys Tyr Met Asn Lys Gln Gln Cys Ile Glu Arg Asp 1 5 10 15

Glu Val Arg Asn Val Phe Arg Glu Leu Glu Ile Leu Gln Glu Ile Glu 20 25 30

His Val Phe Leu Val Asn Leu Trp Tyr Ser Phe Gln Asp Glu Glu Asp 35 40 45

Met Phe Met Val Val Asp Leu Leu Gly Gly Asp Leu Arg Tyr His 50 55 60

| Leu 65 | Gln | Gln | Asn | Val | Gln 70 | Phe | Ser | Glu | Asp | Thr 75 | Val | Arg | Leu | Tyr | Ile 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Cys | Glu | Met | Ala | Leu 85 | Ala | Leu | Asp | Tyr | Leu 90 | Arg | Gly | Gln | His | Ile 95 | Ile |
| His | Arg | Asp | Val 100 | Lys | Pro | Asp | Asn | Ile 105 | Leu | Leu | Asp | Glu | Arg 110 | Gly | His |
| Ala | His | Leu 115 | Thr | Asp | Phe | Asn | Ile 120 | Ala | Thr | Ile | Ile | Lys 125 | Asp | Gly | Glu |
| Arg | Ala 130 | Thr | Ala | Leu | Ala | Gly 135 | Thr | Lys | Pro | Tyr | Met 140 | Ala | Pro | Glu | Ile |
| Phe 145 | His | Ser | Phe | Val | Asn 150 | Gly | Gly | Thr | Gly | Tyr 155 | Ser | Phe | Glu | Val | Asp 160 |
| Trp | Trp | Ser | Val | Gly 165 | Val | Met | Ala | Tyr | Glu 170 | Leu | Leu | Arg | Gly | Trp 175 | Arg |
| Pro | Tyr | Asp | Ile 180 | His | Ser | Ser | Asn | Ala 185 | Val | Glu | Ser | Leu | Val 190 | Gln | Leu |
| Phe | Ser | Thr 195 | Val | Ser | Val | Gln | Tyr 200 | Val | Pro | Thr | Trp | Ser 205 | Lys | Glu | Met |
| Val | Ala 210 | Leu | Leu | Arg | Lys | Leu 215 | Leu | Thr | Val | Asn | Pro 220 | Glu | His | Arg | Leu |
| Ser 225 | Ser | Leu | Gln | Asp | Val 230 | Gln | Ala | Ala | Pro | Ala 235 | Leu | Ala | Gly | Val | Leu 240 |
| Trp | Asp | His | Leu | Ser 245 | Glu | Lys | Arg | Val | Glu 250 | Pro | Gly | Phe | Val | Pro 255 | Asn |
| Lys | Gly | Arg | Leu 260 | His | Cys | Asp | Pro | Thr 265 | Phe | Glu | Leu | Glu | Glu 270 | Met | Ile |
| Leu | Glu | Ser 275 | Arg | Pro | Leu | His | Lys 280 | Lys | Lys | Lys | Arg | Leu 285 | Ala | Lys | Asn |
| Lys | Ser 290 | Arg | Asp | Asn | Ser | Arg 295 | Asp | Ser | Ser | Gln | Ser 300 | Glu | Asn | Asp | Tyr |
| Leu 305 | Gln | Asp | Cys | Leu | Asp 310 | Ala | Ile | Gln | Gln | Asp 315 | Phe | Val | Ile | Phe | Asn 320 |

Ala Pro Glu Ser Arg Asp Ala Ala Glu Pro Val Glu Asp Glu Ala Glu Arg Ser Ala Leu Pro Met Cys Gly Pro Ile Cys Pro Ser Ala Gly Ser Gly <210> 156 <211> 368 <212> PRT <213> Macaca fuscata <400> 156 Met Tyr Ala Met Lys Tyr Met Asn Lys Gln Gln Cys Ile Glu Arg Asp Glu Val Arg Asn Val Phe Arg Glu Leu Gly Ile Leu Gln Glu Ile Glu . His Val Phe Leu Val Asn Leu Trp Tyr Ser Phe Gln Asp Glu Glu Asp Met Phe Met Val Val Asp Leu Leu Gly Gly Asp Leu Arg Tyr His Leu Gln Gln Asn Val Gln Phe Ser Glu Asp Thr Val Arg Leu Tyr Ile Cys Glu Met Ala Leu Ala Leu Asp Tyr Leu Cys Gly Gln His Ile Ile His Arg Asp Val Lys Pro Asp Asn Ile Leu Leu Asp Glu Arg Gly His Ala His Leu Thr Asp Phe Asn Ile Ala Thr Ile Ile Lys Asp Gly Glu Arg Ala Thr Ala Leu Ala Gly Thr Lys Pro Tyr Met Ala Pro Glu Ile

Arg Glu Lys Leu Lys Arg Ser Gln Asp Leu Pro Arg Glu Pro Leu Pro

Phe His Ser Phe Val Asn Gly Gly Thr Gly Tyr Ser Phe Glu Val Asp

Trp Trp Ser Leu Gly Val Met Ala Tyr Glu Leu Leu Arg Gly Trp Arg Pro Tyr Asp Ile His Ser Ser Asn Ala Val Glu Ser Leu Val Gln Leu Phe Ser Thr Val Ser Val Gln Tyr Val Pro Thr Trp Ser Arg Glu Met Val Ala Leu Leu Arg Lys Leu Leu Thr Val Asn Pro Glu His Arg Phe Ser Ser Leu Gln Asp Val Gln Ala Ala Pro Ala Leu Ala Gly Val Leu Trp Gly His Leu Ser Glu Lys Arg Val Glu Pro Asp Phe Val Pro Asn Lys Gly Arg Leu His Cys Asp Pro Thr Phe Glu Leu Glu Glu Met Ile Leu Glu Ser Arg Pro Leu His Lys Lys Lys Lys Arg Leu Ala Lys Asn Lys Ser Arg Asp Asn Ser Arg Asp Ser Ser Gln Ser Glu Asn Asp Tyr Leu Gln Asp Cys Leu Asp Ala Ile Gln Gln Asp Phe Val Ile Phe Asn Arg Glu Lys Leu Lys Arg Ser Gln Asp Leu Pro Ser Glu Pro Leu Pro

<210> 157

<211> 414

<212> PRT

<213> Homo sapiens

Ala Pro Glu Pro Arg Asp Ala Ala Glu Pro Val Glu Asp Glu Glu Gln

Ser Ala Leu Pro Met Cys Gly Pro Ile Cys Pro Ser Ala Gly Ser Gly

| < 400 |)> | 15 |
|-------|----|----|
| Met | G] | У |
| | | |

Gly Asn His Ser His Lys Pro Pro Val Phe Asp Glu Asn Glu

Glu Val Asn Phe Asp His Phe Gln Ile Leu Arg Ala Ile Gly Lys Gly

Ser Phe Gly Lys Val Cys Ile Val Gln Lys Arg Asp Thr Lys Lys Met

Tyr Ala Met Lys Tyr Met Asn Lys Gln Lys Cys Ile Glu Arg Asp Glu

Val Arg Asn Val Phe Arg Glu Leu Gln Ile Met Gln Gly Leu Glu His

Pro Phe Leu Val Asn Leu Trp Tyr Ser Phe Gln Asp Glu Glu Asp Met

Phe Met Val Val Asp Leu Leu Gly Gly Asp Leu Arg Tyr His Leu

Gln Gln Asn Val His Phe Thr Glu Gly Thr Val Lys Leu Tyr Ile Cys

Glu Leu Ala Leu Glu Tyr Leu Gln Arg Tyr His Ile Ile His

Arg Asp Ile Lys Pro Asp Asn Ile Leu Leu Asp Glu His Gly His Val

His Ile Thr Asp Phe Asn Ile Ala Thr Val Val Lys Gly Ala Glu Arg

Ala Ser Ser Met Ala Gly Thr Lys Pro Tyr Met Ala Pro Glu Val Phe

Gln Val Tyr Met Asp Arg Gly Pro Gly Tyr Ser Tyr Pro Val Asp Trp

Trp Ser Leu Gly Ile Thr Ala Tyr Glu Leu Leu Arg Gly Trp Arg Pro

Tyr Glu Ile His Ser Val Thr Pro Ile Asp Glu Ile Leu Asn Met Phe

Lys Val Glu Arg Val His Tyr Ser Ser Thr Trp Cys Lys Gly Met Val

| | | | | 245 | | | | | 250 | | | | | 255 | |
|----------------------------------|--------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala 1 | Leu | Leu | Arg 260 | Lys | Leu | Leu | Thr | Lys 265 | Asp | Pro | Glu | Ser | Arg 270 | Val | Ser |
| Ser 1 | Leu | His 275 | Asp | Ile | Gln | Ser | Val 280 | Pro | Tyr | Leu | Ala | Asp 285 | Met | Asn | Trp |
| Asp A | Ala 290 | Val | Phe | Lys | Lys | Ala 295 | Leu | Met | Pro | Gly | Phe 300 | Val | Pro | Asn | Lys |
| Gly <i>1</i> 305 | Arg | Leu | Asn | Cys | Asp 310 | Pro | Thr | Phe | Glu | Leu 315 | Glu | Glu | Met | Ile | Leu 320 |
| Glu S | Ser | Lys | Pro | Leu 325 | His | Lys | Lys | Lys | Lys 330 | Arg | Leu | Ala | Lys | Asn 335 | Arg |
| Ser A | Arg | Asp | Gly 340 | Thr | Lys | Asp | Ser | Cys 345 | Pro | Leu | Asn | Gly | His 350 | Leu | Gln |
| His (| Cys | Leu 355 | Glu | Thr | Val | Arg | Glu 360 | Glu | Phe | Ile | Ile | Phe 365 | Asn | Arg | Glu |
| Lys 1 | Leu 370 | Arg | Arg | Gln | Gln | Gly 375 | Gln | Gly | Ser | Gln | Leu 380 | Leu | Asp | Thr | Asp |
| Ser <i>1</i> 385 | Arg | Gly | Gly | Gly | Gln 390 | Ala | Gln | Ser | Lys | Leu 395 | Gln | Asp | Gly | Cys | Asn 400 |
| Asn A | Asn | Leu | Leu | Thr 405 | His | Thr | Cys | Thr | Arg 410 | Gly | Cys | Ser | Ser | | |
| <2103 <2113 <2123 <2133 | > 41 > PF | L4 RT | ıscul | lus | | | | | | | | | | | |

| Tyr | Ala 50 | Met | Lys | Tyr | Met | Asn 55 | Lys | Gln | Lys | Cys | Val 60 | Glu | Arg | Asp | Glu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val 65 | Arg | Asn | Val | Phe | Arg 70 | Glu | Leu | Gln | Ile | Met 75 | Gln | Gly | Leu | Glu | His 80 |
| Pro | Phe | Leu | Val | Asn 85 | Leu | Trp | Tyr | Ser | Phe 90 | Gln | Asp | Glu | Glu | Asp 95 | Met |
| Phe | Met | Val | Val 100 | Asp | Leu | Leu | Leu | Gly 105 | Gly | Asp | Leu | Arg | Tyr 110 | His | Leu |
| Gln | Gln | Asn 115 | Val | His | Phe | Thr | Glu 120 | Gly | Thr | Val | Lys | Leu 125 | Tyr | Ile | Cys |
| Glu | Leu 130 | Ala | Leu | Ala | Leu | Glu 135 | Tyr | Leu | Gln | Arg | Tyr 140 | His | Ile | Ile | His |
| Arg 145 | Asp | Ile | Lys | Pro | Asp 150 | Asn | Ile | Leu | Leu | Asp 155 | Glu | His | Gly | His | Val 160 |
| His | Ile | Thr | Asp | Phe 165 | Asn | Ile | Ala | Thr | Val 170 | Leu | Lys | Gly | Ser | Glu 175 | Lys |
| Ala | Ser | Ser | Met 180 | Ala | Gly | Thr | Lys | Pro 185 | Tyr | Met | Ala | Pro | Glu 190 | Val | Phe |
| Gln | Val | Tyr 195 | Val | Asp | Gly | Gly | Pro 200 | Gly | Tyr | Ser | Tyr | Pro 205 | Val | Asp | Trp |
| Trp | Ser 210 | Leu | Gly | Val | Thr | Ala 215 | Tyr | Glu | Leu | Leu | Arg 220 | Gly | Trp | Arg | Pro |
| Tyr 225 | Glu | Ile | His | Ser | Ala 230 | Thr | Pro | Ile | Asp | Glu 235 | Ile | Leu | Asn | Met | Phe 240 |
| Lys | Val | Glu | Arg | Val 245 | His | Tyr | Ser | Ser | Thr 250 | Trp | Cys | Glu | Gly | Met 255 | Val |
| Ser | Leu | Leu | Lys 260 | Lys | Leu | Leu | Thr | Lys 265 | Asp | Pro | Glu | Ser | Arg 270 | Leu | Ser |
| Ser | Leu | Arg 275 | Asp | Ile | Gln | Ser | Met 280 | Thr | Tyr | Leu | Ala | Asp 285 | Met | Asn | Trp |
| Asp | Ala 290 | Val | Phe | Glu | Lys | Ala 295 | Leu | Met | Pro | Gly | Phe | Val | Pro | Asn | Lys |

| 305 | u Asn | Cys | Asp 310 | Pro | Thr | Phe | Glu | Leu 315 | Glu | Glu | Met | Ile | Leu 320 |
|---|------------------------------------|-------------------------------|--------------------------------|-------------------------|--------------------|--------------------------------|--------------------------------|--------------------------------|-------------------------|-------------------------|--------------------------------|-------------------------|--------------------|
| Glu Ser Ly | s Pro | Leu 325 | His | Lys | Lys | Lys | Lys 330 | Arg | Leu | Ala | Lys | His 335 | Arg |
| Ser Arg As | p Ser 340 | Thr | Lys | Asp | Ser | Cys 345 | | Leu | Asn | Gly | His 350 | Leu | Gln |
| Gln Cys Le | | Thr | Val | Arg | Lys 360 | Glu | Phe | Ile | Ile | Phe 365 | Asn | Arg | Glu |
| Lys Leu Ar 370 | g Arg | Gln | Gln | Gly 375 | His | Asp | Gly | Gln | Leu 380 | Ser | Asp | Leu | Asp |
| Gly Arg Il 385 | e Gly | Ser | Gln 390 | Thr | Ser | Ser | Lys | Leu 395 | Gln | Asp | Gly | Arg | Asn 400 |
| Asn Asn Il | e Leu | Thr 405 | His | Thr | Cys | Pro | Arg 410 | Gly | Cys | Ser | Ser | | |
| <210> 159 | | | | | | | | | | | | | |
| <211> 258 <212> PRT | | | | | | | | | | | | | |
| \Z12> EN1 | | | | | | | | | | | | | |
| <213> Homo | sapi | ens | | | | | | | | | | | |
| <213> Homo | _ | | חות ה | Tlo | Clv | Tvo | C1 | Sor | Dho | Clv | Tvo | Vo.1 | Val |
| <213> Homo | _ | | Ala | Ile | Gly | Lys | Gly 10 | Ser | Phe | Gly | Lys | Val 15 | Val |
| <213> Homo | e Leu | Arg 5 | | | | | 10 | | | | | 15 | |
| <213> Homo <400> 159 Phe Gln II 1 Cys Ile Va | e Leu 1 Gln 20 | Arg 5 Lys | Arg | Asp | Thr | Glu 25 | 10 Lys | Met | Tyr | Ala | Met 30 | 15 Lys | Tyr |
| <213> Homo <400> 159 Phe Gln II 1 Cys Ile Va | e Leu 1 Gln 20 s Gln 5 | Arg 5 Lys Gln | Arg Cys | Asp | Thr Glu 40 | Glu 25 Arg | 10 Lys Asp | Met Glu | Tyr Val | Ala Arg 45 | Met 30 Asn | 15 Lys Val | Tyr Phe |
| <213> Homo <400> 159 Phe Gln II 1 Cys Ile Va Met Asn Ly 3 Arg Glu Le | e Leu 1 Gln 20 s Gln 5 | Arg 5 Lys Gln Ile | Arg Cys Leu | Asp Ile Gln 55 | Thr Glu 40 Glu | Glu 25 Arg | 10 Lys Asp Glu | Met Glu His | Tyr Val Val 60 | Ala Arg 45 Phe | Met 30 Asn Leu | 15 Lys Val | Tyr Phe Asn |
| <213> Homo <400> 159 Phe Gln II 1 Cys Ile Va Met Asn Ly 3 Arg Glu Le 50 Leu Trp Ty | e Leu l Gln 20 s Gln 5 u Glu r Ser | Arg 5 Lys Gln Ile | Arg Cys Leu Gln 70 | Asp Ile Gln 55 Asp | Thr Glu 40 Glu Glu | Glu 25 Arg Ile Glu | 10 Lys Asp Glu Asp | Met Glu His Met 75 | Tyr Val Val 60 | Ala Arg 45 Phe | Met 30 Asn Leu Val | 15 Lys Val Val | Tyr Phe Asn Asp 80 |

Leu Asp Tyr Leu Arg Gly Gln His Ile His Arg Asp Val Lys Pro 115 120 125 Asp Asn Ile Leu Leu Asp Glu Arg Gly His Ala His Leu Thr Asp Phe 130 135 140 Asn Ile Ala Thr Ile Ile Lys Asp Gly Glu Arg Ala Thr Ala Leu Ala 145 150 155 Gly Thr Lys Pro Tyr Met Ala Pro Glu Ile Phe His Ser Phe Val Asn 170 175 165 Gly Gly Thr Gly Tyr Ser Phe Glu Val Asp Trp Trp Ser Val Gly Val 185 Met Ala Tyr Glu Leu Leu Arg Gly Trp Arg Pro Tyr Asp Ile His Ser 200 Ser Asn Ala Val Glu Ser Leu Val Gln Leu Phe Ser Thr Val Ser Val 215 220 Gln Tyr Val Pro Thr Trp Ser Lys Glu Met Val Gly Leu Leu Arg Lys 230 235 Val Leu Leu Thr Val Asn Pro Glu His Arg Leu Ser Ser Leu Gln Asp 245 250 255

Val Gln

<210> 160

<211> 252

<212> PRT

<213> Artificial Sequence

<220>

<400> 160

Tyr Glu Leu Glu Val Leu Gly Lys Gly Ala Phe Gly Lys Val Tyr
1 5 10 15

Leu Ala Arg Asp Lys Lys Thr Gly Lys Leu Val Ala Ile Lys Val Ile
20 25 30

Lys Lys Glu Lys Leu Lys Lys Lys Arg Glu Arg Ile Leu Arg Glu Ile Lys Ile Leu Lys Lys Leu Asp His Pro Asn Ile Val Lys Leu Tyr Asp Val Phe Glu Asp Asp Lys Leu Tyr Leu Val Met Glu Tyr Cys Glu Gly Gly Asp Leu Phe Asp Leu Lys Lys Arg Gly Arg Leu Ser Glu Asp Glu Ala Arg Phe Tyr Ala Arg Gln Ile Leu Ser Ala Leu Glu Tyr Leu His Ser Gln Gly Ile Ile His Arg Asp Leu Lys Pro Glu Asn Ile Leu Leu Asp Ser Asp Gly His Val Lys Leu Ala Asp Phe Gly Leu Ala Lys Gln Leu Asp Ser Gly Gly Thr Leu Leu Thr Thr Phe Val Gly Thr Pro Glu Tyr Met Ala Pro Glu Val Leu Leu Gly Lys Gly Tyr Gly Lys Ala Val Asp Ile Trp Ser Leu Gly Val Ile Leu Tyr Glu Leu Leu Thr Gly Lys Pro Pro Phe Pro Gly Asp Asp Gln Leu Leu Ala Leu Phe Lys Lys Ile Gly Lys Pro Pro Pro Pro Pro Pro Pro Pro Glu Trp Lys Ile Ser Pro Glu Ala Lys Asp Leu Ile Lys Lys Leu Leu Val Lys Asp

<210> 161

<211> 255

<212> PRT

<213> Homo sapiens

Pro Glu Lys Arg Leu Thr Ala Glu Glu Ala Leu Glu

| <4 | 00> | 1 | 6 | 1 | L | |
|----|-----|---|---|---|---|--|
| | | _ | | | | |

- Phe Gln Ile Leu Arg Ala Ile Gly Lys Gly Ser Phe Gly Lys Val Val 1 5 10 15
- Cys Ile Val Gln Lys Arg Asp Thr Glu Lys Met Tyr Ala Met Lys Tyr
 20 25 30
- Met Asn Lys Gln Gln Cys Ile Glu Arg Asp Glu Val Arg Asn Val Phe
 35 40 45
- Arg Glu Leu Glu Ile Leu Gln Glu Ile Glu His Val Phe Leu Val Asn 50 55 60
- Leu Trp Tyr Ser Phe Gln Asp Glu Glu Asp Met Phe Met Val Val Asp 65 70 75 80
- Leu Leu Gly Gly Asp Leu Arg Tyr His Leu Gln Gln Asn Val Gln
 85 90 95
- Phe Ser Glu Asp Thr Val Arg Leu Tyr Ile Cys Glu Met Ala Leu Ala 100 105 110
- Leu Asp Tyr Leu Arg Gly Gln His Ile Ile His Arg Asp Val Lys Pro 115 120 125
- Asp Asn Ile Leu Leu Asp Glu Arg Gly His Ala His Leu Thr Asp Phe 130 135 140
- Asn Ile Ala Thr Ile Ile Lys Asp Gly Glu Arg Ala Thr Ala Leu Ala 145 150 155 160
- Gly Thr Lys Pro Tyr Met Ala Pro Glu Ile Phe His Ser Phe Val Asn 165 170 175
- Gly Gly Thr Gly Tyr Ser Phe Glu Val Asp Trp Trp Ser Val Gly Val
 180 185 190
- Met Ala Tyr Glu Leu Leu Arg Gly Trp Arg Pro Tyr Asp Ile His Ser 195 200 205
- Ser Asn Ala Val Glu Ser Leu Val Gln Leu Phe Ser Thr Val Ser Val 210 215 220
- Gln Tyr Val Pro Thr Trp Ser Lys Glu Met Val Gly Leu Leu Arg Lys 225 230 235 240
- Val Leu Leu Thr Val Asn Pro Glu His Arg Leu Ser Ser Leu Gln
 245 250 255

<210> 162 <211> 249 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: pkinase, Protein kinase domain <400> 162 Tyr Glu Leu Gly Glu Lys Leu Gly Ser Gly Ala Phe Gly Lys Val Tyr 10 Lys Gly Lys His Lys Asp Thr Gly Glu Ile Val Ala Ile Lys Ile Leu 20 25 Lys Lys Arg Ser Leu Ser Glu Lys Lys Lys Arg Phe Leu Arg Glu Ile 35 40 45 Gln Ile Leu Arg Arg Leu Ser His Pro Asn Ile Val Arg Leu Leu Gly 50 55 Val Phe Glu Glu Asp Asp His Leu Tyr Leu Val Met Glu Tyr Met Glu 70 75 Gly Gly Asp Leu Phe Asp Tyr Leu Arg Arg Asn Gly Leu Leu Leu Ser 85 90 Glu Lys Glu Ala Lys Lys Ile Ala Leu Gln Ile Leu Arg Gly Leu Glu 105 Tyr Leu His Ser Arg Gly Ile Val His Arg Asp Leu Lys Pro Glu Asn 120 Ile Leu Leu Asp Glu Asn Gly Thr Val Lys Ile Ala Asp Phe Gly Leu 130 135 Ala Arg Lys Leu Glu Ser Ser Ser Tyr Glu Lys Leu Thr Thr Phe Val 145 150 155 160

186

Gly Thr Pro Glu Tyr Met Ala Pro Glu Val Leu Glu Gly Arg Gly Tyr

Ser Ser Lys Val Asp Val Trp Ser Leu Gly Val Ile Leu Tyr Glu Leu

185

170

175

190

165

Leu Thr Gly Lys Leu Pro Phe Pro Gly Ile Asp Pro Leu Glu Glu Leu 195 200 205

Phe Arg Ile Lys Glu Arg Pro Arg Leu Arg Leu Pro Leu Pro Pro Asn 210 215 220

Cys Ser Glu Glu Leu Lys Asp Leu Ile Lys Lys Cys Leu Asn Lys Asp 225 230 235 240

Pro Glu Lys Arg Pro Thr Ala Lys Glu 245

<210> 163

<211> 215

<212> PRT

<213> Homo sapiens

<400> 163

Ile Leu Arg Ala Ile Gly Lys Gly Ser Phe Gly Lys Val Val Cys Ile
1 5 10 15

Val Gln Lys Arg Asp Thr Glu Lys Met Tyr Ala Met Lys Tyr Met Asn 20 25 30

Lys Gln Gln Cys Ile Glu Arg Asp Glu Val Arg Asn Val Phe Arg Glu
35 40 45

Leu Glu Ile Leu Gln Glu Ile Glu His Val Phe Leu Val Asn Leu Trp
50 55 60

Tyr Ser Phe Gln Asp Glu Glu Asp Met Phe Met Val Val Asp Leu Leu 65 70 75 80

Leu Gly Gly Asp Leu Arg Tyr His Leu Gln Gln Asn Val Gln Phe Ser 85 90 95

Glu Asp Thr Val Arg Leu Tyr Ile Cys Glu Met Ala Leu Ala Leu Asp 100 105 110

Tyr Leu Arg Gly Gln His Ile Ile His Arg Asp Val Lys Pro Asp Asn 115 120 125

Ile Leu Leu Asp Glu Arg Gly His Ala His Leu Thr Asp Phe Asn Ile 130 135 140

Ala Thr Ile Ile Lys Asp Gly Glu Arg Ala Thr Ala Leu Ala Gly Thr 145 150 155 160 Lys Pro Tyr Met Ala Pro Glu Ile Phe His Ser Phe Val Asn Gly Gly 165 170 175

Thr Gly Tyr Ser Phe Glu Val Asp Trp Trp Ser Val Gly Val Met Ala 180 185 190

Tyr Glu Leu Leu Arg Gly Trp Arg Pro Tyr Asp Ile His Ser Ser Asn 195 200 205

Ala Val Glu Ser Leu Val Gln 210 215

<210> 164

<211> 216

<212> PRT

<213> Artificial Sequence

<220>

<400> 164

Leu Gly Lys Lys Leu Gly Glu Gly Ala Phe Gly Glu Val Tyr Lys Gly
1 5 10 15

Thr Leu Lys Gly Lys Gly Gly Val Glu Val Glu Val Ala Val Lys Thr
20 25 30

Leu Lys Glu Asp Ala Ser Glu Gln Gln Ile Glu Glu Phe Leu Arg Glu 35 40 45

Ala Arg Leu Met Arg Lys Leu Asp His Pro Asn Ile Val Lys Leu Leu 50 55 60

Gly Val Cys Thr Glu Glu Glu Pro Leu Met Ile Val Met Glu Tyr Met 65 70 75 80

Glu Gly Gly Asp Leu Leu Asp Tyr Leu Arg Lys Asn Arg Pro Lys Glu 85 90 95

Leu Ser Leu Ser Asp Leu Leu Ser Phe Ala Leu Gln Ile Ala Arg Gly
100 105 110

Met Glu Tyr Leu Glu Ser Lys Asn Phe Val His Arg Asp Leu Ala Ala 115 120 125 Arg Asn Cys Leu Val Gly Glu Asn Lys Thr Val Lys Ile Ala Asp Phe Gly Leu Ala Arg Asp Leu Tyr Asp Asp Tyr Tyr Arg Lys Lys Ser Pro Arg Leu Pro Ile Arg Trp Met Ala Pro Glu Ser Leu Lys Asp Gly Lys Phe Thr Ser Lys Ser Asp Val Trp Ser Phe Gly Val Leu Leu Trp Glu Ile Phe Thr Leu Gly Glu Ser Pro Tyr Pro Gly Met Ser Asn Glu Glu Val Leu Glu Tyr Leu Lys <210> 165 <211> 187 <212> PRT <213> Homo sapiens <400> 165 Met Gln Cys Leu Leu Thr Leu Ser Met Ala Leu Val Cys Ala Ile Gln Ala Arg Asp Ile Pro Gln Thr Lys Gln Asp Val Glu Leu Pro Lys Leu Ala Gly Thr Trp Tyr Ser Met Ala Met Val Ala Ser Asp Phe Ser Leu Leu Glu Thr Val Glu Ala Pro Leu Arg Val Asn Ile Thr Ser Leu Trp Pro Thr Pro Glu Gly Asn Leu Glu Ile Ile Leu His Arg Trp Glu His His Arg Cys Val Glu Arg Thr Val Leu Ala Gln Lys Thr Glu Asp Pro Ala Val Phe Met Val Asp Arg Ile Cys Arg Ala Ala Val Val

Ser Gly Gln Gln Pro Ser Gln Arg Trp Arg Leu Ser Val Lys Glu Arg

Ser Arg Lys Glu Gly Gly Arg Leu Pro Arg Ser Arg Asp Lys Lys Asp 130 135 140

Leu Cys Val Gly His Arg Leu Asp Asp Arg Ser Tyr Val Phe Phe Cys
145 150 155 160

Met Gly Thr Thr Pro Ser Ala Asp His His Thr Met Cys Gln Tyr 165 170 175

Leu Gly Met Thr Gln Gly Pro Pro Gly Phe Ile 180 185

<210> 166

<211> 186

<212> PRT

<213> PAPIO CYNOCEPHALUS

<400> 166

Met Gln Cys Leu Leu Thr Leu Gly Val Ala Leu Ile Cys Gly Val
1 5 10 15

Trp Ala Ile Asn Ser Pro Gln Thr Met Gln Asp Val Glu Leu Pro Lys
20 25 30

Leu Ala Gly Thr Trp His Ser Met Ala Met Ala Ala Ser Asp Phe Ser 35 40 45

Leu Leu Glu Thr Lys Glu Ala Pro Leu Arg Ile Tyr Ile Ser Ser Leu 50 55 60

Gln Pro Thr Pro Glu Gly Asn Leu Glu Ile Val Leu Arg Arg Trp Ser 65 70 75 80

Gln Lys Gln Ser Pro Phe Arg Asp Ser Asn Gln Cys Ile Glu Glu Lys 85 90 95

Ile Ile Ala Glu Lys Thr Glu Asn Pro Ile Glu Phe Lys Ile Asn Tyr 100 105 110

Leu Asp Glu Asn Arg Ile Tyr Leu Phe Asn Thr Asp Gly Ser Lys Tyr
115 120 125

Leu Phe Leu Cys Leu Glu Ser Thr Arg Arg Gln Asn Leu Ala Cys Gln 130 135 140

Tyr Leu Ala Arg Thr Leu Glu Val Asp Asp Lys Val Met Ala Glu Phe

145 150 155 160

Ile Ser Phe Leu Lys Thr Leu Pro Val His Met Gln Ile Phe Leu Asp 165 170 175

Met Thr Gln Ala Glu Glu Gln Cys Arg Val 180 185

<210> 167

<211> 180

<212> PRT

<213> Homo sapiens

<400> 167

Met Leu Cys Leu Leu Thr Leu Gly Val Ala Leu Val Cys Gly Val
1 5 10 . 15

Pro Ala Met Asp Ile Pro Gln Thr Lys Gln Asp Leu Glu Leu Pro Lys
20 25 30

Leu Ala Gly Thr Trp His Ser Met Ala Met Ala Thr Asn Asn Ile Ser
35 40 45

Leu Met Ala Thr Leu Lys Ala Pro Leu Arg Val His Ile Thr Ser Leu 50 55 60

Leu Pro Thr Pro Glu Asp Asn Leu Glu Ile Val Leu His Arg Trp Glu 65 70 75 80

Asn Asn Ser Cys Val Glu Lys Lys Val Leu Gly Glu Lys Thr Glu Asn 85 90 95

Pro Lys Lys Phe Lys Ile Asn Tyr Thr Val Ala Asn Glu Ala Thr Leu 100 105 110

Leu Asp Thr Asp Tyr Asp Asn Phe Leu Phe Leu Cys Leu Gln Asp Thr 115 120 125

Thr Thr Pro Ile Gln Ser Met Met Cys Gln Tyr Leu Ala Arg Val Leu 130 135 140

Val Glu Asp Asp Glu Ile Met Gln Gly Phe Ile Arg Ala Phe Arg Pro 145 150 155 160

Leu Pro Arg His Leu Trp Tyr Leu Leu Asp Leu Lys Gln Met Glu Glu 165 170 175 Pro Cys Arg Phe 180

<210> 168

<211> 188

<212> PRT

<213> Homo sapiens

<400> 168

Ser Glu Pro Pro Thr Ala Ala Ala Met Leu Cys Leu Leu Leu Thr Leu 1 5 10 15

Gly Val Ala Leu Val Cys Gly Val Pro Ala Met Asp Ile Pro Gln Thr
20 25 30

Lys Gln Asp Leu Glu Leu Pro Lys Leu Ala Gly Thr Trp His Ser Met 35 40 45

Ala Met Ala Thr Asn Asn Ile Ser Leu Met Ala Thr Leu Lys Ala Pro 50 55 60

Leu Arg Val His Ile Thr Ser Leu Leu Pro Thr Pro Glu Asp Asn Leu 65 70 75 80

Glu Ile Val Leu His Arg Trp Glu Asn Asn Ser Cys Val Glu Lys Lys
85 90 95

Val Leu Gly Glu Lys Thr Glu Asn Pro Lys Lys Phe Lys Ile Asn Tyr
100 105 110

Thr Val Ala Asn Glu Ala Thr Leu Leu Asp Thr Asp Tyr Asp Asn Phe 115 120 125

Leu Phe Leu Cys Leu Gln Asp Thr Thr Thr Pro Ile Gln Ser Met Met 130 135 140

Cys Gln Tyr Leu Ala Arg Val Leu Val Glu Asp Asp Glu Ile Met Gln 145 150 155 160

Gly Phe Ile Arg Ala Phe Arg Pro Leu Pro Arg His Leu Trp Tyr Leu 165 170 175

Leu Asp Leu Lys Gln Met Glu Glu Pro Cys Arg Phe 180 185

<210> 169

<211> 163 <212> PRT <213> Felis catus <400> 169 Ala Thr Leu Pro Pro Thr Met Glu Asp Leu Asp Ile Arg Gln Val Ala Gly Thr Trp His Ser Met Ala Met Ala Ala Ser Asp Ile Ser Leu Leu Asp Ser Glu Thr Ala Pro Leu Arg Val Tyr Val Gln Glu Leu Arg Pro Thr Pro Arg Asp Asn Leu Glu Ile Ile Leu Arg Lys Arg Glu Asn His Ala Cys Ile Glu Gly Asn Ile Met Ala Gln Arg Thr Glu Asp Pro Ala Val Phe Met Val Asp Tyr Gln Gly Glu Lys Lys Ile Ser Val Leu Asp Thr Asp Tyr Thr His Tyr Met Phe Phe Cys Met Glu Ala Pro Ala Pro Gly Thr Glu Asn Gly Met Met Cys Gln Tyr Leu Ala Arg Thr Leu Lys Ala Asp Asn Glu Val Met Glu Lys Phe Asp Arg Ala Leu Gln Thr Leu Pro Val His Ile Arg Ile Ile Leu Asp Leu Thr Gln Gly Lys Glu Gln Cys Arg Val <210> 170 <211> 145 <212> PRT <213> Homo sapiens

Lys Phe Ala Gly Lys Trp Tyr Leu Val Ala Ser Ala Asn Phe Asp Pro

<400> 170

Thr Pro Leu Lys Glu Gly Asn Leu Glu Ile Val Phe Asp Gly Asp Lys 35 40 45 Asn Gly Ile Cys Glu Glu Thr Phe Gly Lys Leu Glu Lys Thr Lys Lys 50 55 60 Leu Gly Val Glu Phe Asp Tyr Tyr Thr Gly Asp Asn Arg Phe Val Val 65 70 75 80 Leu Asp Thr Asp Tyr Asp Asn Tyr Leu Leu Val Cys Val Gln Lys Gly 85 90 Asp Gly Asn Glu Thr Ser Arg Thr Ala Glu Leu Tyr Gly Arg Thr Pro 100 105 Glu Leu Ser Pro Glu Ala Leu Glu Leu Phe Glu Thr Ala Thr Lys Glu 115 120 125 Leu Gly Ile Pro Glu Asp Asn Val Val Cys Thr Arg Gln Thr Glu Arg 130 135 140 Cys 145 <210> 171 <211> 145 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: lipocalin domain sequence <400> 171 Lys Phe Ala Gly Lys Trp Tyr Leu Val Ala Ser Ala Asn Phe Asp Pro 10 Glu Leu Lys Glu Glu Leu Gly Val Leu Glu Ala Thr Arg Lys Glu Ile 30 20 25 Thr Pro Leu Lys Glu Gly Asn Leu Glu Ile Val Phe Asp Gly Asp Lys Asn Gly Ile Cys Glu Glu Thr Phe Gly Lys Leu Glu Lys Thr Lys Lys

Glu Leu Lys Glu Glu Leu Gly Val Leu Glu Ala Thr Arg Lys Glu Ile

25

30

50 55 60

Leu Gly Val Glu Phe Asp Tyr Tyr Thr Gly Asp Asn Arg Phe Val Val 65 70 75 80

Leu Asp Thr Asp Tyr Asp Asn Tyr Leu Leu Val Cys Val Gln Lys Gly 85 90 95

Asp Gly Asn Glu Thr Ser Arg Thr Ala Glu Leu Tyr Gly Arg Thr Pro 100 105 110

Glu Leu Ser Pro Glu Ala Leu Glu Leu Phe Glu Thr Ala Thr Lys Glu 115 120 125

Leu Gly Ile Pro Glu Asp Asn Val Val Cys Thr Arg Gln Thr Glu Arg, 130 135 140

Cys 145

<210> 172

<211> 1327

<212> PRT

<213> Mus musculus

<400> 172

Met Glu Ala Pro Leu Gln Thr Gly Met Val Leu Gly Val Met Ile Gly 1 5 10 15

Ala Gly Val Ala Val Leu Val Thr Ala Val Leu Ile Leu Leu Val Val 20 25 30

Arg Arg Leu Arg Val Gln Lys Thr Pro Ala Pro Glu Gly Pro Arg Tyr 35 40 45

Arg Phe Arg Lys Arg Asp Lys Val Leu Phe Tyr Gly Arg Lys Ile Met 50 55 60

Arg Lys Val Ser Gln Ser Thr Ser Ser Leu Val Asp Thr Ser Val Ser 65 70 75 80

Thr Thr Ser Arg Pro Arg Met Lys Lys Leu Lys Met Leu Asn Ile
85 90 95

Ala Lys Lys Ile Leu Arg Ile Gln Lys Glu Thr Pro Thr Leu Gln Arg
100 105 110

Lys Glu Pro Pro Pro Ser Val Leu Glu Ala Asp Leu Thr Glu Gly Asp Leu Ala Asn Ser His Leu Pro Ser Glu Val Leu Tyr Met Leu Lys Asn Val Arg Val Leu Gly His Phe Glu Lys Pro Leu Phe Leu Glu Leu Cys Arg His Met Val Phe Gln Arg Leu Gly Gln Gly Asp Tyr Val Phe Arg Pro Gly Gln Pro Asp Ala Ser Ile Tyr Val Val Gln Asp Gly Leu Leu Glu Leu Cys Leu Pro Gly Pro Asp Gly Lys Glu Cys Val Val Lys Lys Val Val Pro Gly Asp Ser Val Asn Ser Leu Leu Ser Ile Leu Asp Val Ile Thr Gly His Gln His Pro Gln Arg Thr Val Ser Ala Arg Ala Ala Arg Asp Ser Thr Val Leu Arg Leu Pro Val Glu Ala Phe Ser Ala Val Phe Thr Lys Tyr Pro Glu Ser Leu Val Arg Val Val Gln Ile Ile Met Val Arg Leu Gln Arg Val Thr Phe Leu Ala Leu His Asn Tyr Leu Gly Leu Thr Asn Glu Leu Phe Ser His Glu Ile Gln Pro Leu Arg Leu Phe Pro Ser Pro Gly Leu Pro Thr Arg Thr Ser Pro Val Arg Gly Ser Lys Arg Val Val Ser Thr Ser Gly Thr Glu Asp Thr Ser Lys Glu Thr Ser Gly Arg Pro Leu Asp Ser Ile Gly Ala Pro Leu Pro Gly Pro Ala Gly Asp Pro Val Lys Pro Thr Ser Leu Glu Ala Pro Pro Ala Pro Leu Leu

Ser Arg Cys Ile Ser Met Pro Val Asp Ile Ser Gly Leu Gln Gly Gly Pro Arg Ser Asp Phe Asp Met Ala Tyr Glu Arg Gly Arg Ile Ser Val Ser Leu Gln Glu Glu Ala Ser Gly Gly Pro Gln Thr Ala Ser Pro Arg Thr Pro Thr Gln Glu Leu Arg Glu Gln Pro Ala Gly Ala Cys Glu Tyr Ser Tyr Cys Glu Asp Glu Ser Ala Thr Gly Gly Cys Pro Phe Gly Pro Tyr Gln Gly Arg Gln Thr Ser Ser Ile Phe Glu Ala Ala Lys Arg Glu Leu Ala Lys Leu Met Arg Ile Glu Asp Pro Ser Leu Leu Asn Ser Arg Val Leu Leu His His Ala Lys Ala Gly Thr Ile Ile Ala Arg Gln Gly Asp Gln Asp Val Ser Leu His Phe Val Leu Trp Gly Cys Leu His Val Tyr Gln Arg Met Ile Asp Lys Ala Glu Glu Val Cys Leu Phe Val Ala Gln Pro Gly Glu Leu Val Gly Gln Leu Ala Val Leu Thr Gly Glu Pro Leu Ile Phe Thr Leu Arg Ala Gln Arg Asp Cys Thr Phe Leu Arg Ile Ser Lys Ser His Phe Tyr Glu Ile Met Arg Ala Gln Pro Ser Val Val Leu Ser Ala Ala His Thr Val Ala Ala Arg Met Ser Pro Phe Val Arg Gln Met Asp Phe Ala Ile Asp Trp Thr Ala Val Glu Ala Gly Arg Ala Leu Tyr Arg Gln Gly Asp Arg Ser Asp Cys Thr Tyr Ile Val Leu Asn

| Gly 625 | Arg | Leu | Arg | Ser | Val 630 | Ile | Gln | Arg | Gly | Ser 635 | Gly | Lys | Lys | Glu | Leu 640 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val | Gly | Glu | Tyr | Gly 645 | Arg | Gly | Asp | Leu | Ile 650 | Gly | Val | Val | Glu | Ala 655 | Leu |
| Thr | Arg | Gln | Pro 660 | Arg | Ala | Thr | Thr | Val 665 | His | Ala | Val | Arg | Asp 670 | Thr | Glu |
| Leu | Ala | Lys 675 | Leu | Pro | Glu | Gly | Thr 680 | Leu | Gly | His | Ile | Lys 685 | Arg | Arg | Tyr |
| Pro | Gln 690 | Val | Val | Thr | Arg | Leu 695 | Ile | His | Leu | Leu | Ser 700 | Gln | Lys | Ile | Leu |
| Gly 705 | Asn | Leu | Gln | Gln | Leu 710 | Gln | Gly | Pro | Phe | Pro 715 | Gly | Ser | Gly | Leu | Ser 720 |
| Val | Pro | Gln | His | Ser 725 | Glu | Leu | Thr | Asn | Pro 730 | Ala | Ser | Asn | Leu | Ser 735 | Thr |
| Val | Ala | Ile | Leu 740 | Pro | Val | Cys | Ala | Glu 745 | Val | Pro | Met | Met | Ala 750 | Phe | Thr |
| Leu | Glu | Leu 755 | Gln | His | Ala | Leu | Gln 760 | Ala | Ile | Gly | Pro | Thr 765 | Leu | Leu | Leu |
| Asn | Ser 770 | Asp | Val | Ile | Arg | Ala 775 | Leu | Leu | Gly | Ala | Ser 780 | Ala | Leu | Asp | Ser |
| Ile 785 | Gln | Glu | Phe | Arg | Leu 790 | Ser | Gly | Trp | Leu | Ala 795 | Gln | Gln | Glu | Asp | Ala 800 |
| His | Arg | Ile | Val | Leu 805 | Tyr | Gln | Thr | Asp | Thr 810 | Ser | Leu | Thr | Pro | Trp 815 | Thr |
| Val | Arg | Cys | Leu 820 | Arg | Gln | Ala | Asp | Cys 825 | Ile | Leu | Ile | Val | Gly 830 | Leu | Gly |
| Asp | Gln | Glu 835 | Pro | Thr | Val | Gly | Gln 840 | Leu | Glu | Gln | Met | Leu 845 | Glu | Asn | Thr |
| Ala | Val 850 | Arg | Ala | Leu | Lys | Gln 855 | Leu | Val | Leu | Leu | His 860 | Arg | Glu | Glu | Gly |
| Pro 865 | Gly | Pro | Thr | Arg | Thr 870 | Val | Glu | Trp | Leu | Asn 875 | Met | Arg | Ser | Trp | Cys 880 |

Ser Gly His Leu His Leu Arg Cys Pro Arg Arg Leu Phe Ser Arg Arg Ser Pro Ala Lys Leu His Glu Leu Tyr Glu Lys Val Phe Ser Arg Arg Ala Asp Arg His Ser Asp Phe Ser Arg Leu Ala Arg Val Leu Thr Gly Asn Thr Ile Ala Leu Val Leu Gly Gly Gly Ala Arg Gly Cys Ser His Ile Gly Val Leu Lys Ala Leu Glu Glu Ala Gly Val Pro Val Asp Leu Val Gly Gly Thr Ser Ile Gly Ser Phe Ile Gly Ala Leu Tyr Ala Glu Glu Arg Ser Ala Ser Arg Thr Lys Gln Arg Ala Arg Glu Trp Ala Lys Ser Met Thr Ser Val Leu Glu Pro Val Leu Asp Leu Thr Tyr Pro Val Thr Ser Met Phe Thr Gly Ser Ala Phe Asn Arg Ser Ile His Arg Val Phe Gln Asp Lys Gln Ile Glu Asp Leu Trp Leu Pro Tyr Phe Asn Val Thr Thr Asp Ile Thr Ala Ser Ala Met Arg Val His Lys Asp Gly Ser Leu Trp Arg Tyr Val Arg Ala Ser Met Thr Leu Ser Gly Tyr Leu Pro Pro Leu Cys Asp Pro Lys Asp Gly His Leu Leu Met Asp Gly Gly Tyr Ile Asn Asn Leu Pro Ala Asp Ile Ala Arg Ser Met Gly Ala Lys Thr Val Ile Ala Ile Asp Val Gly Ser Gln Asp Glu Thr Asp Leu Ser Thr Tyr Gly Asp Ser Leu Ser Gly Trp Trp Leu Leu Trp Lys Arg Leu

Asn Pro Trp Ala Asp Lys Val Lys Val Pro Asp Met Ala Glu Ile Gln Ser Arg Leu Ala Tyr Val Ser Cys Val Arg Gln Leu Glu Val Val Lys Ser Ser Ser Tyr Cys Glu Tyr Leu Arg Pro Ser Ile Asp Cys Phe Lys Thr Met Asp Phe Gly Lys Phe Asp Gln Ile Tyr Asp Val Gly Tyr Gln Tyr Gly Lys Ala Val Phe Gly Gly Trp Thr Arg Gly Glu Val Ile Glu Lys Met Leu Thr Asp Arg Arg Ser Thr Asp Leu Asn Glu Ser Arg Arg Ala Asp Ile Leu Ala Phe Pro Ser Ser Gly Phe Thr Asp Leu Ala Glu Ile Val Ser Arg Ile Glu Pro Pro Thr Ser Tyr Val Ser Asp Gly Cys Ala Asp Gly Glu Glu Ser Asp Cys Leu Thr Glu Tyr Glu Glu Asp Ala Gly Pro Asp Cys Ser Arg Asp Glu Gly Gly Ser Pro Glu Gly Ala Ser Pro Ser Thr Ala Ser Glu Val Glu Glu Lys Ser Thr Leu Arg Gln Arg Arg Phe Leu Pro Gln Glu Thr Pro Ser Ser Val Ala Asp Ala <210> 173 <211> 702 <212> PRT <213> Homo sapiens <400> 173 Met Leu Ser Gly Arg Leu Arg Ser Val Ile Arg Lys Asp Asp Gly Lys

Lys Arg Leu Ala Gly Glu Tyr Gly Arg Gly Asp Leu Val Gly Val Val

| Glu | Thr | Leu 35 | Thr | His | Gln | Ala | Arg 40 | Ala | Thr | Thr | Val | His 45 | Ala | Val | Arg |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Asp | Ser 50 | Glu | Leu | Ala | Lys | Leu 55 | Pro | Ala | Gly | Ala | Leu 60 | Thr | Ser | Ile | Lys |
| Arg 65 | Arg | Tyr | Pro | Gln | Val 70 | Val | Thr | Arg | Leu | Ile 75 | His | Leu | Leu | Gly | Glu 80 |
| Lys | Ile | Leu | Gly | Ser 85 | Leu | Gln | Gln | Gly | Pro 90 | Val | Thr | Gly | His | Gln 95 | Leu |
| Gly | Leu | Pro | Thr 100 | Glu | Gly | Ser | Lys | Trp 105 | Asp | Leu | Gly | Asn | Pro 110 | Ala | Val |
| Asn | Leu | Ser 115 | Thr | Val | Ala | Val | Met 120 | Pro | Val | Ser | Glu | Glu 125 | Val | Pro | Leu |
| Thr | Ala 130 | Phe | Ala | Leu | Glu | Leu 135 | Glu | His | Ala | Leu | Ser 140 | Ala | Ile | Gly | Pro |
| Thr 145 | Leu | Leu | Leu | Thr | Ser 150 | Asp | Asn | Ile | Lys | Arg 155 | Arg | Leu | Gly | Ser | Ala 160 |
| Ala | Leu | Asp | Ser | Val 165 | His | Glu | Tyr | Arg | Leu 170 | Ser | Ser | Trp | Leu | Gly 175 | Gln |
| Gln | Glu | Asp | Thr 180 | His | Arg | Ile | Val | Leu 185 | Tyr | Gln | Ala | Asp | Gly 190 | Thr | Leu |
| Thr | Pro | Trp 195 | Thr | Gln | Arg | Cys | Val 200 | Arg | Gln | Ala | Asp | Cys 205 | Ile | Leu | Ile |
| Val | Gly 210 | Leu | Gly | Asp | Gln | Glu 215 | Pro | Thr | Val | Gly | Glu 220 | Leu | Glu | Arg | Met |
| Leu 225 | Glu | Ser | Thr | Ala | Val 230 | Arg | Ala | Gln | Lys | Gln 235 | Leu | Ile | Leu | Leu | His 240 |
| Arg | Glu | Glu | Gly | Pro 245 | Ala | Pro | Ala | Arg | Thr 250 | Val | Glu | Trp | Leu | Asn 255 | Met |
| Arg | Ser | Ser | Cys 260 | Ser | Gly | His | Leu | His 265 | Leu | Cys | Cys | Pro | Arg 270 | Arg | Val |
| Phe | Ser | Arg | Arg | Ser | Leu | Pro | Lys | Leu | Val | Glu | Met | Tyr | Lys | His | Val |

| Phe | Gln 290 | Arg | Pro | Pro | Asp | Arg 295 | His | Ser | Asp | Phe | Ser 300 | Arg | Leu | Ala | Arg |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val 305 | Leu | Thr | Gly | Asn | Ala 310 | Ile | Ala | Leu | Val | Leu 315 | Gly | Gly | Gly | Gly | Ala 320 |
| Arg | Gly | Cys | Ala | Gln 325 | Val | Gly | Val | Leu | Lys 330 | Ala | Leu | Ala | Glu | Cys 335 | Gly |
| Ile | Pro | Val | Asp 340 | Met | Val | Gly | Gly | Thr 345 | Ser | Ile | Gly | Ala | Phe 350 | Val | Gly |
| Ala | Leu | Tyr 355 | Ser | Glu | Glu | Arg | Asn 360 | Tyr | Ser | Gln | Met | Arg 365 | Ile | Arg | Ala |
| Lys | Gln 370 | Trp | Ala | Glu | Gly | Met 375 | Thr | Ser | Leu | Met | Lys 380 | Ala | Ala | Leu | Asp |
| Leu 385 | Thr | Tyr | Pro | Ile | Thr 390 | Ser | Met | Phe | Ser | Gly 395 | Ala | Gly | Phe | Asn | Ser 400 |
| Ser | Ile | Phe | Ser | Val 405 | Phe | Lys | Asp | Gln | Gln 410 | Ile | Glu | Asp | Leu | Trp 415 | Ile |
| Pro | Tyr | Phe | Ala 420 | Ile | Thr | Thr | Asp | Ile 425 | Thr | Ala | Ser | Ala | Met 430 | Arg | Val |
| His | Thr | Asp 435 | Gly | Ser | Leu | Trp | Arg 440 | Tyr | Val | Arg | Ala | Ser 445 | Met | Ser | Leu |
| Ser | Gly 450 | Tyr | Met | Pro | Pro | Leu 455 | Cys | Asp | Pro | Lys | Asp 460 | Gly | His | Leu | Leu |
| Met 465 | Asp | Gly | Gly | Tyr | Ile 470 | Asn | Asn | Leu | Pro | Ala 475 | Asp | Val | Ala | Arg | Ser 480 |
| Met | Gly | Ala | Lys | Val 485 | Val | Ile | Ala | Ile | Asp 490 | Val | Gly | Ser | Arg | Asp 495 | Glu |
| Thr | Asp | Leu | Thr 500 | Asn | Tyr | Gly | Asp | Ala 505 | Leu | Ser | Gly | Trp | Trp 510 | Leu | Leu |
| Trp | Lys | Arg 515 | Trp | Asn | Pro | Leu | Ala 520 | Thr | Lys | Val | Lys | Val 525 | Leu | Asn | Met |
| Ala | Glu 530 | Ile | Gln | Thr | Arg | Leu 535 | Ala | Tyr | Val | Cys | Cys 540 | Val | Arg | Gln | Leu |

Glu Val Val Lys Ser Ser Asp Tyr Cys Glu Tyr Leu Arg Pro Pro Ile 545 550 555 Asp Ser Tyr Ser Thr Leu Asp Phe Gly Lys Phe Asn Glu Ile Cys Glu 565 570 575 Val Gly Tyr Gln His Gly Arg Thr Val Phe Asp Ile Trp Gly Arg Ser 580 585 Gly Val Leu Glu Lys Met Leu Arg Asp Gln Gln Gly Pro Ser Lys Lys 600 Pro Ala Ser Ala Val Leu Thr Cys Pro Asn Ala Ser Phe Thr Asp Leu 615 Ala Glu Ile Val Ser Arg Ile Glu Pro Ala Lys Pro Ala Met Val Asp 630 635 Asp Glu Ser Asp Tyr Gln Thr Glu Tyr Glu Glu Leu Leu Asp Val 645 650 Pro Arg Asp Ala Tyr Ala Asp Phe Gln Ser Thr Ser Ala Gln Gln Gly 665 Ser Asp Leu Glu Asp Glu Ser Ser Leu Arg His Arg His Pro Ser Leu 675 680 685 Ala Phe Pro Lys Leu Ser Glu Gly Ser Ser Asp Gln Asp Gly 690 695 700 <210> 174 <211> 1425 <212> PRT <213> Drosophila melanogaster <400> 174 Met Asp Val Leu Glu Met Leu Arg Ala Ser Ala Ser Gly Ser Tyr Asn Thr Thr Phe Ser Asp Ala Trp Cys Gln Tyr Val Ser Lys Gln Ile Thr 20 25 Ala Thr Val Tyr Met Tyr Phe Ala Leu Val Met Met Ser Leu Leu Phe 45

Ile Ala Trp Phe Leu Tyr Phe Lys Arg Met Ala Arg Leu Arg Leu Arg

| Asp | Glu | Ile | Ala | Arg | Ser | Ile | Ser | Thr | Val | Thr | Asn | Ser | Ser | Gly | Asp |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |

60

55

50

75 75

Met Arg Gly Leu Arg Phe Arg Lys Arg Asp Lys Met Leu Phe Tyr Gly 85 90 95

Arg Arg Met Leu Arg Lys Met Lys Asn Val Ser Gly Gln Met Tyr Ser 100 105 110

Ser Gly Lys Gly Tyr Lys Arg Arg Ala Val Met Arg Phe Ala Arg Arg 115 120 125

Ile Leu Gln Leu Arg Arg Asp Asn Met Pro Leu Glu Met Arg Thr Val 130 135 140

Glu Pro Pro Ala Glu Tyr Leu Glu Glu Thr Ile Glu Gly Ser Asp Arg 145 150 155 160

Val Pro Pro Asp Ala Leu Tyr Met Leu Gln Ser Ile Arg Ile Phe Gly
165 170 175

His Phe Glu Lys Pro Val Phe Leu Arg Leu Cys Lys His Thr Gln Leu 180 185 190

Leu Glu Leu Met Ala Gly Asp Tyr Leu Phe Lys Ile Thr Asp Pro Asp 195 200 205

Asp Ser Val Tyr Ile Val Gln Ser Gly Met Ile Asn Val Tyr Ile Ser 210 215 220

Asn Ala Asp Gly Ser Thr Leu Ser Leu Lys Thr Val Arg Lys Gly Glu 225 230 235 240

Ser Val Thr Ser Leu Leu Ser Phe Ile Asp Val Leu Ser Gly Asn Pro 245 250 255

Ser Tyr Tyr Lys Thr Val Thr Ala Lys Ala Ile Glu Lys Ser Val Val 260 265 270

Ile Arg Leu Pro Met Gln Ala Phe Glu Glu Val Phe Gln Asp Asn Pro 275 280 285

Asp Val Met Ile Arg Val Ile Gln Val Ile Met Ile Arg Leu Gln Arg 290 295 300

Val Leu Phe Thr Ala Leu Arg Asn Tyr Leu Gly Leu Asn Ala Glu Leu

| 305 | | 310 | | | 315 | | | 320 |
|--------------------|------------------|----------------|--------------|----------------|----------------|----------------|------------|------------|
| Val Gln Asn | His Met 2 325 | Arg Tyr | Lys S | Ser Val 330 | Ser Thr | Met Ser | Gly 335 | Pro |
| Ile Asn Ser | Gln Thr S | Ser Gln | | Ser Arg 345 | Gln Ala | Pro Asn 350 | Gly | Pro |
| Pro Met Val | Ile Ser (| Gln Met | Asn I 360 | Leu Met | Gln Ser | Ala Val 365 | Ser | Gly |
| Thr Gly Ser 370 | Ser Gly | Val Ser 375 | Val T | Thr Val | Thr Arg 380 | Pro Pro | Ser | Ser |
| Pro Ser Arg 385 | | Arg Glu 390 | Glu H | His Thr | Leu Ser 395 | Asp Pro | Asn | Pro 400 |
| Asn Pro Asp | Gly Ser 3 | Phe His | Gly T | Thr Thr 410 | Asn Leu | Phe Thr | Glu 415 | Val |
| His Gly Asp | Ala Pro A | Asn Ala | - | Leu Phe 425 | His Gln | Gln Gln 430 | Gln | Gln |
| His Ser Val 435 | Gly Asn | Leu Ser | Thr A | Arg Arg | Ser Ser | Ile Thr 445 | Leu | Met |
| Ala Pro Asp 450 | Pro Ser | His Ser 455 | Cys I | Leu Gln | Thr Pro | Gly Val | Thr | Thr |
| Ser Ile Asp 465 | | Leu Val 470 | Gln S | Ser Ser | Ala Val 475 | Asp Ser | Leu | Arg 480 |
| Lys Glu Leu | Gly Leu 8 | Ser Glu | Glu A | Asp Ser 490 | His Ile | Ile Glu | Pro 495 | Phe |
| Val Glu Leu | Arg Glu : | Leu Glu | | Asn Val | Thr Leu | Ile Thr 510 | Glu | Gly |
| Asn Ala Asp 515 | Asp Val | Cys Val | Trp E | Phe Val | Met Thr | Gly Thr 525 | Leu | Ala |
| Val Tyr Gln 530 | Ser Asn | Gln Asp 535 | Ala T | Thr Arg | Ala Lys 540 | Gln Asp | Lys | Ser |

Asp Met Leu Ile His Phe Val His Pro Gly Glu Ile Val Gly Gly Leu

Ala Met Leu Thr Gly Glu Ala Ser Ala Tyr Thr Ile Arg Ser Arg Ser

- Ile Thr Arg Ile Ala Phe Ile Arg Arg Ala Ala Ile Tyr Gln Ile Met 580 585 590
- Arg Gln Arg Pro Arg Ile Val Leu Asp Leu Gly Asn Gly Val Val Arg 595 600 605
- Arg Leu Ser Pro Leu Val Arg Gln Cys Asp Tyr Ala Leu Asp Trp Ile 610 620
- Phe Leu Glu Ser Gly Arg Ala Val Tyr Arg Gln Asp Glu Ser Ser Asp 625 630 635 640
- Ser Thr Tyr Ile Val Leu Ser Gly Arg Met Arg Ser Val Ile Thr His 645 650 655
- Pro Gly Gly Lys Lys Glu Ile Val Gly Glu Tyr Gly Lys Gly Asp Leu 660 665 670
- Val Gly Ile Val Glu Met Ile Thr Glu Thr Ser Arg Thr Thr Thr Val 675 680 685
- Met Ala Val Arg Asp Ser Glu Leu Ala Lys Leu Pro Glu Gly Leu Phe 690 695 700
- Asn Ala Ile Lys Leu Arg Tyr Pro Ile Val Val Thr Lys Leu Ile Ser 705 710 715 720
- Phe Leu Ser His Arg Phe Leu Gly Ser Met Gln Thr Arg Ser Gly Ser 725 730 735
- Gly Ala Pro Gly Ala Pro Val Glu Ala Asn Pro Val Thr His Lys Tyr
 740 745 750
- Ser Thr Val Ala Leu Val Pro Ile Thr Asp Glu Val Pro Met Thr Pro
 755 760 765
- Phe Thr Tyr Glu Leu Tyr His Ser Leu Cys Ala Ile Gly Pro Val Leu 770 775 780
- His Leu Thr Ser Asp Val Val Arg Lys Gln Leu Gly Ser Asn Ile Phe 785 790 795 800
- Glu Ala Ala Asn Glu Tyr Arg Leu Thr Ser Trp Leu Ala Gln Glu 805 810 815
- Asp Arq Asn Ile Ile Thr Leu Tyr Gln Cys Asp Ser Ser Leu Ser Ala

820

Trp Thr Gln Arg Cys Met Arg Gln Ala Asp Val Ile Leu Ile Val Gly 835 840 845

825

- Leu Gly Asp Arg Ser His Leu Val Gly Lys Phe Glu Arg Glu Ile Asp 850 855 860
- Arg Leu Ala Met Arg Thr Gln Lys Glu Leu Val Leu Leu Tyr Pro Glu 865 870 875 880
- Ala Ser Asn Ala Lys Pro Ala Asn Thr Leu Ser Trp Leu Asn Ala Arg 885 890 895
- Pro Trp Val Thr Lys His His His Val Leu Cys Val Lys Arg Ile Phe 900 905 910
- Thr Arg Lys Ser Gln Tyr Arg Ile Asn Asp Leu Tyr Ser Arg Val Leu 915 920 925
- Leu Ser Glu Pro Asn Met His Ser Asp Phe Ser Arg Leu Ala Arg Trp 930 935 940
- Leu Thr Gly Asn Ser Ile Gly Leu Val Leu Gly Gly Gly Gly Ala Arg 945 950 955 960
- Gly Ala Ala His Ile Gly Met Leu Lys Ala Ile Gln Glu Ala Gly Ile 965 970 975
- Pro Val Asp Met Val Gly Gly Val Ser Ile Gly Ala Leu Met Gly Ala 980 985 990
- Leu Trp Cys Ser Glu Arg Asn Ile Thr Thr Val Thr Gln Lys Ala Arg 995 1000 1005
- Glu Trp Ser Lys Lys Met Thr Lys Trp Phe Leu Gln Leu Leu Asp Leu 1010 1015 1020
- Thr Tyr Pro Ile Thr Ser Met Phe Ser Gly Arg Glu Phe Asn Lys Thr 1025 1030 1035 1040
- Ile His Asp Thr Phe Gly Asp Val Ser Ile Glu Asp Leu Trp Ile Pro 1045 1050 1055
- Tyr Phe Thr Leu Thr Thr Asp Ile Thr Ala Ser Cys His Arg Ile His
 1060 1065 1070
- Thr Asn Gly Ser Leu Trp Arg Tyr Val Arg Ser Ser Met Ser Leu Ser

| 1075 1080 1 | .08 |
|-------------|-----|
|-------------|-----|

- Gly Tyr Met Pro Pro Leu Cys Asp Pro Lys Asp Gly His Leu Leu Leu 1090 1095 1100
- Asp Gly Gly Tyr Val Asn Asn Leu Pro Ala Asp Val Met His Asn Leu 1105 1110 1115 1120
- Gly Ala Ala His Ile Ile Ala Ile Asp Val Gly Ser Gln Asp Asp Thr 1125 1130 1135
- Asp Leu Thr Asn Tyr Gly Asp Asp Leu Ser Gly Trp Trp Leu Leu Tyr 1140 1150
- Lys Lys Trp Asn Pro Phe Thr Ser Pro Val Lys Val Pro Asp Leu Pro 1155 1160 1165
- Asp Ile Gln Ser Arg Leu Ala Tyr Val Ser Cys Val Arg Gln Leu Glu 1170 1175 1180
- Glu Val Lys Asn Ser Asp Tyr Cys Glu Tyr Ile Arg Pro Pro Ile Asp 1185 1190 1195 1200
- Lys Tyr Lys Thr Leu Ala Phe Gly Ser Phe Asp Glu Ile Arg Asp Val 1205 1210 1215
- Gly Tyr Val Phe Gly Lys Asn Tyr Phe Glu Ser Met Ala Lys Ala Gly 1220 1230
- Arg Leu Gly Arg Phe Asn Gln Trp Phe Asn Lys Glu Pro Pro Lys Arg 1235 1240 1245
- Val Asn His Ala Ser Leu Asn Glu Tyr Thr Phe Ile Asp Leu Ala Gln 1250 1255 1260
- Ile Val Cys Arg Leu Pro Glu Thr Tyr Ala Val Asn Thr Ala Glu Leu 1265 1270 1275 1280
- Phe Ser Glu Asp Glu Asp Cys Asp Gly Tyr Ile Ser Glu Pro Thr Thr
 1285 1290 1295
- Leu Asn Thr Asp Arg Arg Ile Gln Val Ser Arg Ala Gly Asn Ser 1300 1305 1310
- Leu Ser Phe Ser Glu Thr Glu Met Asp Ser Asp Val Glu Leu Asp Leu 1315 1320 1325
- Lys Leu Glu Arg Lys Thr Asp Lys Ser Thr Gln Ser Ser Pro Pro Ser

| 1330 | 1335 | 1340 |
|------|------|------|
| | | |

Asn Ser Arg Ser Asp Met Arg Gly Lys Glu Glu Ala Arg His Met Ser 1345 1350 1355 1360

Asn Trp His Trp Gly Val Lys His Lys Asp Glu Thr Gly Ser Gly Ala 1365 1370 1375

Asn Glu Ala Thr Lys Thr Gln Thr Gly Gln Glu Gln Glu Leu Gln Gln 1380 1385 1390

Glu Gln Gln Asp Gln Gly Ala Thr Ala Glu Gln Leu Val Asp Lys Asp 1395 1400 1405

Lys Glu Glu Asn Lys Glu Asn Arg Ser Ser Pro Asn Asn Glu Thr Lys 1410 1415 1420

Asn 1425

<210> 175

<211> 1389

<212> PRT

<213> Drosophila melanogaster

<400> 175

Met Tyr Phe Ala Leu Val Met Met Ser Leu Leu Phe Ile Ala Trp Phe 1 5 10 15

Leu Tyr Phe Lys Arg Met Ala Arg Leu Arg Leu Arg Asp Glu Ile Ala 20 25 30

Arg Ser Ile Ser Thr Val Thr Asn Ser Ser Gly Asp Met Arg Gly Leu
35 40 45

Arg Phe Arg Lys Arg Asp Lys Met Leu Phe Tyr Gly Arg Arg Met Leu 50 55 60

Arg Lys Met Lys Asn Val Ser Gly Gln Met Tyr Ser Ser Gly Lys Gly 65 70 75 80

Tyr Lys Arg Arg Ala Val Met Arg Phe Ala Arg Arg Ile Leu Gln Leu 85 90 95

Arg Arg Asp Asn Met Pro Leu Glu Met Arg Thr Val Glu Pro Pro Ala 100 105 110

| Glu | Tyr | Leu 115 | Glu | Glu | Thr | Ile | Glu 120 | Gly | Ser | Asp | Arg | Val 125 | Pro | Pro | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Leu 130 | Tyr | Met | Leu | Gln | Ser 135 | Ile | Arg | Ile | Phe | Gly 140 | His | Phe | Glu | Lys |
| Pro 145 | Val | Phe | Leu | Arg | Leu 150 | Суѕ | Lys | His | Thr | Gln 155 | Leu | Leu | Glu | Leu | Met 160 |
| Ala | Gly | Asp | Tyr | Leu 165 | Phe | Lys | Ile | Thr | Asp 170 | Pro | Asp | Asp | Ser | Val 175 | Tyr |
| Ile | Val | Gln | Ser 180 | Gly | Met | Ile | Asn | Val 185 | Tyr | Ile | Ser | Asn | Ala 190 | Asp | Gly |
| Ser | Thr | Leu 195 | Ser | Leu | Lys | Thr | Val 200 | Arg | Lys | Gly | Glu | Ser 205 | Val | Thr | Ser |
| Leu | Leu 210 | Ser | Phe | Ile | Asp | Val 215 | Leu | Ser | Gly | Asn | Pro 220 | Ser | Tyr | Tyr | Lys |
| Thr 225 | Val | Thr | Ala | Lys | Ala 230 | Ile | Glu | Lys | Ser | Val 235 | Val | Ile | Arg | Leu | Pro 240 |
| Met | Gln | Ala | Phe | Glu 245 | Glu | Val | Phe | Gln | Asp 250 | Asn | Pro | Asp | Val | Met 255 | Ile |
| Arg | Val | Ile | Gln 260 | Val | Ile | Met | Ile | Arg 265 | Leu | Gln | Arg | Val | Leu 270 | Phe | Thr |
| Ala | Leu | Arg 275 | Asn | Tyr | Leu | Gly | Leu 280 | Asn | Ala | Glu | Leu | Val 285 | Gln | Asn | His |
| Met | Arg 290 | Tyr | Lys | Ser | Val | Ser 295 | Thr | Met | Ser | Gly | Pro 300 | Ile | Asn | Ser | Gln |
| Thr 305 | Ser | Gln | Ser | Ser | Arg 310 | Gln | Ala | Pro | Asn | Gly 315 | Pro | Pro | Met | Val | Ile 320 |
| Ser | Gln | Met | Asn | Leu 325 | Met | Gln | Ser | Ala | Val 330 | Ser | Gly | Thr | Gly | Ser 335 | Ser |
| Gly | Val | Ser | Val 340 | Thr | Val | Thr | Arg | Pro 345 | Pro | Ser | Ser | Pro | Ser 350 | Arg | His |
| Ser | Arg | Glu 355 | Glu | His | Thr | Leu | Ser 360 | Asp | Pro | Asn | Pro | Asn 365 | Pro | Asp | Gly |

| Ser | Phe 370 | His | Gly | Thr | Thr | Asn 375 | Leu | Phe | Thr | Glu | Val 380 | His | Gly | Asp | Ala |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Pro 385 | Asn | Ala | Asp | Leu | Phe 390 | His | Gln | Gln | Gln | Gln 395 | Gln | His | Ser | Val | Gly 400 |
| Asn | Leu | Ser | Thr | Arg 405 | Arg | Ser | Ser | Ile | Thr 410 | Leu | Met | Ala | Pro | Asp 415 | Gly |
| Ser | His | Ser | Cys 420 | Leu | Gln | Thr | Pro | Gly 425 | Val | Thr | Thr | Ser | Ile 430 | Asp | Met |
| Arg | Leu | Val 435 | Gln | Ser | Ser | Ala | Val 440 | Asp | Ser | Leu | Arg | Lys 445 | Glu | Leu | Gly |
| Leu | Ser 450 | Glu | Glu | Asp | Ser | His 455 | Ile | Ile | Glu | Pro | Phe 460 | Val | Glu | Leu | Arg |
| Glu 465 | Leu | Glu | Pro | Asn | Val 470 | Thr | Leu | Ile | Thr | Glu 475 | Gly | Asn | Ala | Asp | Asp 480 |
| Val | Cys | Val | Trp | Phe 485 | Val | Met | Thr | Gly | Thr 490 | Leu | Ala | Val | Tyr | Gln 495 | Ser |
| Asn | Gln | Asp | Ala 500 | Thr | Arg | Ala | Lys | Gln 505 | Asp | Lys | Ser | Asp | Met 510 | Leu | Ile |
| His | Phe | Val 515 | His | Pro | Gly | Glu | Ile 520 | Val | Gly | Gly | Leu | Ala 525 | Met | Leu | Thr |
| Gly | Glu 530 | Ala | Ser | Ala | Tyr | Thr 535 | Ile | Arg | Ser | Arg | Ser 540 | Ile | Thr | Arg | Ile |
| Ala 545 | Phe | Ile | Arg | Arg | Ala 550 | Ala | Ile | Tyr | Gln | Ile 555 | Met | Arg | Gln | Arg | Pro 560 |
| Arg | Ile | Val | Leu | Asp 565 | Leu | Gly | Asn | Gly | Val 570 | Val | Arg | Arg | Leu | Ser 575 | Pro |
| Leu | Val | Arg | Gln 580 | Cys | Asp | Tyr | Ala | Leu 585 | Asp | Trp | Ile | Phe | Leu 590 | Glu | Ser |
| Gly | Arg | Ala 595 | Val | Tyr | Arg | Gln | Asp 600 | Glu | Ser | Ser | Asp | Ser 605 | Thr | Tyr | Ile |
| Val | Leu 610 | Ser | Gly | Arg | Met | Arg 615 | Ser | Val | Ile | Thr | His 620 | Pro | Gly | Gly | Lys |

| Lys Glu II 625 | le Val Gl | y Glu T 630 | yr Gly | Lys | Gly | Asp 635 | Leu | Val | Gly | Ile | Val 640 |
|-------------------|------------------|-----------------|---------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu Met I | le Thr Gl 64 | | er Arg | Thr | Thr 650 | Thr | Val | Met | Ala | Val 655 | Arg |
| Asp Ser G | lu Leu Al 660 | a Lys L | eu Pro | Glu 665 | Gly | Leu | Phe | Asn | Ala 670 | Ile | Lys |
| Leu Arg Ty | yr Pro Il 75 | e Val V | al Thr 680 | Lys | Leu | Ile | Ser | Phe 685 | Leu | Ser | His |
| Arg Phe Le | eu Gly Se | | ln Thr 95 | Arg | Ser | Gly | Ser 700 | Gly | Ala | Pro | Gly |
| Ala Pro Va 705 | al Glu Al | a Asn P. 710 | ro Val | Thr | His | Lys 715 | Tyr | Ser | Thr | Val | Ala 720 |
| Leu Val P | co Ile Th 72 | _ | lu Val | Pro | Met 730 | Thr | Pro | Phe | Thr | Tyr 735 | Glu |
| Leu Tyr H | is Ser Le 740 | u Cys A | la Ile | Gly 745 | Pro | Val | Leu | Arg | Leu 750 | Thr | Ser |
| Asp Val V | al Arg Ly 55 | s Gln L | eu Gly 760 | Ser | Asn | Ile | Phe | Glu 765 | Ala | Ala | Asn |
| Glu Tyr A | rg Leu Th | | rp Leu 75 | Ala | Gln | Gln | Glu 780 | Asp | Arg | Asn | Ile |
| Ile Thr Lo | eu Tyr Gl | n Cys A 790 | sp Ser | Ser | Leu | Ser 795 | Ala | Trp | Thr | Gln | Arg 800 |
| Cys Met A | rg Gln Al 80 | | al Ile | Leu | Ile 810 | Val | Gly | Leu | Gly | Asp 815 | Arg |
| Ser His L | eu Val Gl 820 | y Lys P | he Glu | Arg 825 | Glu | Ile | Asp | Arg | Leu 830 | Ala | Met |
| Arg Thr G | ln Lys Gl 35 | u Leu V | al Leu 840 | Leu | Tyr | Pro | Glu | Ala 845 | Ser | Asn | Ala |
| Lys Pro A | la Asn Th | | er Trp 55 | Leu | Asn | Ala | Arg 860 | Pro | Trp | Val | Thr |
| Lys His H: 865 | is His Va | l Leu C 870 | ys Val | Lys | Arg | Ile 875 | Phe | Thr | Arg | Lys | Ser 880 |

| Gln Tyr Arg | Ile Asn 885 | Asp Leu | Tyr Ser | Arg Val | . Leu | Leu | Ser | Glu 895 | Pro |
|-----------------------|--------------------|-----------------|-----------------|-----------------|-------------|-------------|-------------|-------------|-------------|
| Asn Met His | Ser Asp 900 | Phe Ser | Arg Leu 905 | Ala Arg | J Trp | Leu | Thr 910 | Gly | Asn |
| Ser Ile Gly 915 | Leu Val | Leu Gly | Gly Gly 920 | Gly Ala | a Arg | Gly 925 | Ala | Ala | His |
| Ile Gly Met 930 | Leu Lys | Ala Ile 935 | Gln Glu | Ala Gly | 940 | Pro | Val | Asp | Met |
| Val Gly Gly 945 | Val Ser | Ile Gly 950 | Ala Leu | Met Gly | | Leu | Trp | Cys | Ser 960 |
| Glu Arg Asn | Ile Thr 965 | Thr Val | Thr Gln | Lys Ala 970 | a Arg | Glu | Trp | Ser 975 | Lys |
| Lys Met Thr | Lys Trp 980 | Phe Leu | Gln Leu 985 | Leu Ası | Leu | Thr | Туг 990 | Pro | Ile |
| Thr Ser Met 995 | Phe Ser | | Glu Phe 1000 | Asn Lys | | Ile 1005 | His | Asp | Thr |
| Phe Gly Asp 1010 | Val Ser | Ile Glu 1015 | Asp Leu | Trp Ile | Pro 1020 | Tyr | Phe | Thr | Leu |
| Thr Thr Asp | | Ala Ser .030 | Cys His | Arg Ile 103 | | Thr | Asn | | Ser L040 |
| Leu Trp Arg | Tyr Val 1045 | Arg Ser | Ser Met | Ser Lei 1050 | ı Ser | Gly | _ | Met L055 | Pro |
| Pro Leu Cys 1 | Asp Pro 060 | Lys Asp | Gly His | | ı Leu | _ | Gly 1070 | Gly | Tyr |
| Val Asn Asn | | | | | | | 77- | 7/1 = | Иic |
| 1075 | Leu Pro | | Val Met 1080 | His Ası | | Gly 1085 | Ата | AIa | 1113 |
| 1075 Ile Ile Ala 1090 | | : | 1080 | | | 1085 | | | |
| Ile Ile Ala | Ile Asp Asp Leu | Val Gly 1095 | 1080 Ser Gln | Asp Asp | Thr 1100 | 1085 Asp | Leu | Thr | Asn |

- Arg Leu Ala Tyr Val Ser Cys Val Arg Gln Leu Glu Glu Val Lys Asn 1140 1145 1150
- Ser Asp Tyr Cys Glu Tyr Ile Arg Pro Pro Ile Asp Lys Tyr Lys Thr 1155 1160 1165
- Leu Ala Phe Gly Ser Phe Asp Glu Ile Arg Asp Val Gly Tyr Val Phe 1170 1175 1180
- Gly Lys Asn Tyr Phe Glu Ser Met Ala Lys Ala Gly Arg Leu Gly Arg 1185 1190 1195 1200
- Phe Asn Gln Trp Phe Asn Lys Glu Pro Pro Lys Arg Val Asn His Ala 1205 1210 1215
- Ser Leu Asn Glu Tyr Thr Phe Ile Asp Leu Ala Gln Ile Val Cys Arg 1220 1225 1230
- Leu Pro Glu Thr Tyr Ala Val Asn Thr Ala Glu Leu Phe Ser Glu Asp 1235 1240 1245
- Glu Asp Cys Asp Gly Tyr Ile Ser Glu Pro Thr Thr Leu Asn Thr Asp 1250 1255 1260
- Arg Arg Ile Gln Val Ser Arg Ala Gly Asn Ser Leu Ser Phe Ser 1265 1270 1275 1280
- Glu Thr Glu Met Asp Ser Asp Val Glu Leu Asp Leu Lys Leu Glu Arg 1285 1290 1295
- Lys Thr Asp Lys Ser Thr Gln Ser Ser Pro Pro Ser Asn Ser Arg Ser 1300 1305 1310
- Asp Met Arg Gly Lys Glu Glu Ala Arg His Met Ser Asn Trp His Trp 1315 1320 1325
- Gly Val Lys His Lys Asp Glu Thr Gly Ser Gly Ala Thr Glu Ala Thr 1330 1335 1340
- Lys Thr Gln Thr Gly Gln Glu Gln Glu Leu Gln Gln Gln Gln Gln Asp 1345 1350 1355 1360
- Gln Gly Ala Thr Ala Glu Gln Leu Val Asp Lys Asp Lys Glu Glu Asn 1365 1370 1375
- Lys Glu Asn Arg Ser Ser Pro Asn Asn Glu Thr Lys Asn 1380 1385

<210> 176 <211> 1327 <212> PRT <213> Homo sapiens <400> 176 Met Glu Ala Pro Leu Gln Thr Gly Met Val Leu Gly Val Met Ile Gly Ala Gly Val Ala Val Val Thr Ala Val Leu Ile Leu Leu Val Val Arg Arg Leu Arg Val Pro Lys Thr Pro Ala Pro Asp Gly Pro Arg Tyr Arg Phe Arg Lys Arg Asp Lys Val Leu Phe Tyr Gly Arg Lys Ile Met Arg Lys Val Ser Gln Ser Thr Ser Ser Leu Val Asp Thr Ser Val Ser Ala Thr Ser Arg Pro Arg Met Arg Lys Lys Leu Lys Met Leu Asn Ile Ala Lys Lys Ile Leu Arg Ile Gln Lys Glu Thr Pro Thr Leu Gln Arg Lys Glu Pro Pro Pro Ala Val Leu Glu Ala Asp Leu Thr Glu Gly Asp Leu Ala Asn Ser His Leu Pro Ser Glu Val Leu Tyr Met Leu Lys Asn Val Arg Val Leu Gly His Phe Glu Lys Pro Leu Phe Leu Glu Leu Cys Arg His Met Val Phe Gln Arg Leu Gly Gln Gly Asp Tyr Val Phe Arg Pro Gly Gln Pro Asp Ala Ser Ile Tyr Val Val Gln Asp Gly Leu Leu Glu Leu Cys Leu Pro Gly Pro Asp Gly Lys Glu Cys Val Val Lys Glu

Val Val Pro Gly Asp Ser Val Asn Ser Leu Leu Ser Ile Leu Asp Val

| Ile 225 | Thr | Gly | His | Gln | His 230 | Pro | Gln | Arg | Thr | Val 235 | Ser | Ala | Arg | Ala | Ala 240 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Arg | Asp | Ser | Thr | Val 245 | Leu | Arg | Leu | Pro | Val 250 | Glu | Ala | Phe | Ser | Ala 255 | Val |
| Phe | Thr | Lys | Tyr 260 | Pro | Glu | Ser | Leu | Val 265 | Arg | Val | Val | Gln | Ile 270 | Ile | Met |
| Val | Arg | Leu 275 | Gln | Arg | Val | Thr | Phe 280 | Leu | Ala | Leu | His | Asn 285 | Tyr | Leu | Gly |
| Leu | Thr 290 | Asn | Glu | Leu | Phe | Ser 295 | His | Glu | Ile | Gln | Pro 300 | Leu | Arg | Leu | Phe |
| Pro 305 | Ser | Pro | Gly | Leu | Pro 310 | Thr | Arg | Thr | Ser | Pro 315 | Val | Arg | Gly | Ser | Lys 320 |
| Arg | Met | Val | Ser | Thr 325 | Ser | Ala | Thr | Asp | Glu 330 | Pro | Arg | Glu | Thr | Pro 335 | Gly |
| Arg | Pro | Pro | Asp 340 | Pro | Thr | Gly | Ala | Pro 345 | Leu | Pro | Gly | Pro | Thr 350 | Gly | Asp |
| Pro | Val | Lys 355 | Pro | Thr | Ser | Leu | Glu 360 | Thr | Pro | Ser | Ala | Pro 365 | Leu | Leu | Ser |
| Arg | Cys 370 | Val | Ser | Met | Pro | Gly 375 | Asp | Ile | Ser | Gly | Leu 380 | Gln | Gly | Gly | Pro |
| Arg 385 | Ser | Asp | Phe | Asp | Met 390 | Ala | Tyr | Glu | Arg | Gly 395 | Arg | Ile | Ser | Val | Ser 400 |
| Leu | Gln | Glu | Glu | Ala 405 | Ser | Gly | Gly | Ser | Leu 410 | Ala | Ala | Pro | Ala | Arg 415 | Thr |
| Pro | Thr | Gln | Glu 420 | Pro | Arg | Glu | Gln | Pro 425 | Ala | Gly | Ala | Cys | Glu 430 | Tyr | Ser |
| Tyr | Cys | Glu 435 | Asp | Glu | Ser | Ala | Thr 440 | Gly | Gly | Cys | Pro | Phe 445 | Gly | Pro | Tyr |
| Gln | Gly 450 | Arg | Gln | Thr | Ser | Ser 455 | Ile | Phe | Glu | Ala | Ala 460 | Lys | Gln | Glu | Leu |
| Ala 465 | Lys | Leu | Met | Arg | Ile 470 | Glu | Asp | Pro | Ser | Leu 475 | Leu | Asn | Ser | Arg | Val 480 |

| Leu | Leu | His | His | Ala 485 | Lys | Ala | Gly | Thr | Ile 490 | Ile | Ala | Arg | Gln | Gly 495 | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gln | Asp | Val | Ser 500 | Leu | His | Phe | Val | Leu 505 | Trp | Gly | Cys | Leu | His 510 | Val | Tyr |
| Gln | Arg | Met 515 | Ile | Asp | Lys | Ala | Glu 520 | Asp | Val | Cys | Leu | Phe 525 | Val | Ala | Gln |
| Pro | Gly 530 | Glu | Leu | Val | Gly | Gln 535 | Leu | Ala | Val | Leu | Thr 540 | Gly | Glu | Pro | Leu |
| Ile 545 | Phe | Thr | Leu | Arg | Ala 550 | Gln | Arg | Asp | Cys | Thr 555 | Phe | Leu | Arg | Ile | Ser 560 |
| Lys | Ser | Asp | Phe | Tyr 565 | Glu | Ile | Met | Arg | Ala 570 | Gln | Pro | Ser | Val | Val 575 | Leu |
| Ser | Ala | Ala | His 580 | Thr | Val | Ala | Ala | Arg 585 | Met | Ser | Pro | Phe | Val 590 | Arg | Gln |
| Met | Asp | Phe 595 | Ala | Ile | Asp | Trp | Thr 600 | Ala | Val | Glu | Ala | Gly 605 | Arg | Ala | Leu |
| Tyr | Arg 610 | Gln | Gly | Asp | Arg | Ser 615 | Asp | Cys | Thr | Tyr | Ile 620 | Val | Leu | Asn | Gly |
| Arg 625 | Leu | Arg | Ser | Val | Ile 630 | Gln | Arg | Gly | Ser | Gly 635 | Lys | Lys | Glu | Leu | Val 640 |
| Gly | Glu | Tyr | Gly | Arg 645 | Gly | Asp | Leu | Ile | Gly 650 | Val | Val | Glu | Ala | Leu 655 | Thr |
| Arg | Gln | Pro | Arg 660 | Ala | Thr | Thr | Val | His 665 | Ala | Val | Arg | Asp | Thr 670 | Glu | Leu |
| Ala | Lys | Leu 675 | Pro | Glu | Gly | Thr | Leu 680 | Gly | His | Ile | Lys | Arg 685 | Arg | Tyr | Pro |
| Gln | Val 690 | Val | Thr | Arg | Leu | Ile 695 | His | Leu | Leu | Ser | Gln 700 | Lys | Ile | Leu | Gly |
| Asn 705 | Leu | Gln | Gln | Leu | Gln 710 | Gly | Pro | Phe | Pro | Ala 715 | Gly | Ser | Gly | Leu | Gly 720 |
| Val | Pro | Pro | His | Ser 725 | Glu | Leu | Thr | Asn | Pro 730 | Ala | Ser | Asn | Leu | Ala 735 | Thr |

Val Ala Ile Leu Pro Val Cys Ala Glu Val Pro Met Val Ala Phe Thr Leu Glu Leu Gln His Ala Leu Gln Ala Ile Gly Pro Thr Leu Leu Leu Asn Ser Asp Ile Ile Arg Ala Arg Leu Gly Ala Ser Ala Leu Asp Ser Ile Gln Glu Phe Arg Leu Ser Gly Trp Leu Ala Gln Gln Glu Asp Ala His Arg Ile Val Leu Tyr Gln Thr Asp Ala Ser Leu Thr Pro Trp Thr Val Arg Cys Leu Arg Gln Ala Asp Cys Ile Leu Ile Val Gly Leu Gly Asp Gln Glu Pro Thr Leu Gly Gln Leu Glu Gln Met Leu Glu Asn Thr Ala Val Arg Ala Leu Lys Gln Leu Val Leu Leu His Arg Glu Glu Gly Ala Gly Pro Thr Arg Thr Val Glu Trp Leu Asn Met Arg Ser Trp Cys Ser Gly His Leu His Leu Arg Cys Pro Arg Arg Leu Phe Ser Arg Arg Ser Pro Ala Lys Leu His Glu Leu Tyr Glu Lys Val Phe Ser Arg Arg Ala Asp Arg His Ser Asp Phe Ser Arg Leu Ala Arg Val Leu Thr Gly Asn Thr Ile Ala Leu Val Leu Gly Gly Gly Gly Ala Arg Gly Cys Ser His Ile Gly Val Leu Lys Ala Leu Glu Glu Ala Gly Val Pro Val Asp Leu Val Gly Gly Thr Ser Ile Gly Ser Phe Ile Gly Ala Leu Tyr Ala Glu Glu Arg Ser Ala Ser Arg Thr Lys Gln Arg Ala Arg Glu Trp Ala

- Lys Ser Met Thr Ser Val Leu Glu Pro Val Leu Asp Leu Thr Tyr Pro 995 1000 1005
- Val Thr Ser Met Phe Thr Gly Ser Ala Phe Asn Arg Ser Ile His Arg 1010 1015 1020
- Val Phe Gln Asp Lys Gln Ile Glu Asp Leu Trp Leu Pro Tyr Phe Asn 1025 1030 1035 1040
- Val Thr Thr Asp Ile Thr Ala Ser Ala Met Arg Val His Lys Asp Gly 1045 1050 1055
- Ser Leu Trp Arg Tyr Val Arg Ala Ser Met Thr Leu Ser Gly Tyr Leu 1060 1065 1070
- Pro Pro Leu Cys Asp Pro Lys Asp Gly His Leu Leu Met Asp Gly Gly
 1075 1080 1085
- Tyr Ile Asn Asn Leu Pro Ala Asp Ile Ala Arg Ser Met Gly Ala Lys 1090 1095 1100
- Thr Val Ile Ala Ile Asp Val Gly Ser Gln Asp Glu Thr Asp Leu Ser 1105 1110 1115 1120
- Thr Tyr Gly Asp Ser Leu Ser Gly Trp Trp Leu Leu Trp Lys Arg Leu 1125 1130 1135
- Asn Pro Trp Ala Asp Lys Val Lys Val Pro Asp Met Ala Glu Ile Gln
 1140 1145 1150
- Ser Arg Leu Ala Tyr Val Ser Cys Val Arg Gln Leu Glu Val Val Lys 1155 1160 1165
- Ser Ser Ser Tyr Cys Glu Tyr Leu Arg Pro Pro Ile Asp Cys Phe Lys 1170 1175 1180
- Thr Met Asp Phe Gly Lys Phe Asp Gln Ile Tyr Asp Val Gly Tyr Gln 1185 1190 1195 1200
- Tyr Gly Lys Ala Val Phe Gly Gly Trp Ser Arg Gly Asn Val Ile Glu 1205 1210 1215
- Lys Met Leu Thr Asp Arg Arg Ser Thr Asp Leu Asn Glu Ser Arg Arg 1220 1225 1230
- Ala Asp Val Leu Ala Phe Pro Ser Ser Gly Phe Thr Asp Leu Ala Glu 1235 1240 1245

Ile Val Ser Arg Ile Glu Pro Pro Thr Ser Tyr Val Ser Asp Gly Cys 1250 1255 1260

Ala Asp Gly Glu Glu Ser Asp Cys Leu Thr Glu Tyr Glu Glu Asp Ala 1265 1270 1275 1280

Gly Pro Asp Cys Ser Arg Asp Glu Gly Gly Ser Pro Glu Gly Ala Ser 1285 1290 1295

Pro Ser Thr Ala Ser Glu Met Glu Glu Glu Lys Ser Ile Leu Arg Gln 1300 1305 1310

Arg Arg Cys Leu Pro Gln Glu Pro Pro Gly Ser Ala Thr Asp Ala 1315 1320 1325

<210> 177

<211> 331

<212> PRT

<213> Homo sapiens

<400> 177

Pro Asp Arg His Ser Asp Phe Ser Arg Leu Ala Arg Val Leu Thr Gly
1 5 10 15

Asn Ala Ile Ala Leu Val Leu Gly Gly Gly Gly Ala Ser Met Thr Ser 20 25 30

Leu Met Lys Ala Ala Leu Asp Leu Thr Tyr Pro Ile Thr Ser Met Phe
35 40 45

Ser Gly Ala Gly Phe Asn Ser Ser Ile Phe Ser Val Phe Lys Asp Gln 50 55 60

Gln Ile Glu Asp Leu Trp Ile Pro Tyr Phe Ala Ile Thr Thr Asp Ile
65 70 75 80

Thr Ala Ser Ala Met Arg Val His Thr Asp Gly Ser Leu Trp Trp Tyr 85 90 95

Val Arg Ala Ser Met Ser Leu Ser Gly Tyr Met Pro Pro Leu Cys Asp 100 105 110

Pro Lys Asp Gly His Leu Leu Met Asp Gly Gly Tyr Ile Asn Asn Leu 115 120 125

Pro Ala Ala Ser Ala Pro Arg Ser Leu Gly Trp Asn Thr Phe Ser Leu

Glu Tyr Ala Lys Gly Lys Cys Gln Ala Gly Ile Arg Ala Pro Arg Thr
145 150 155 160

Cys Thr Arg Val Tyr Met His Thr Gln Ala Pro Ala Ala Cys Ala Pro 165 170 175

Ala Tyr Gly Pro Val Cys Gln Leu Ser Ser Met Gln Asn Lys Gly Gln
180 185 190

Val Glu Glu Leu Gly Ala Ile Lys Pro His Leu Cys Pro Gln Ser Glu 195 200 205

Thr Asn Ser Leu Gln Gly Val Thr Arg Ala Gly Phe Ser Leu Ala Asp 210 215 220

Val Ala Arg Ser Met Gly Ala Lys Val Val Ile Ala Ile Asp Val Gly 225 230 235 240

Ser Arg Asp Glu Thr Asp Leu Thr Asn Tyr Gly Asp Ala Leu Ser Gly 245 250 255

Trp Trp Leu Leu Trp Lys Arg Trp Asn Pro Leu Ala Thr Lys Val Lys 260 265 270

Val Leu Asn Met Ala Glu Ile Gln Thr Arg Leu Ala Tyr Val Cys Cys 275 280 285

Val Arg Gln Leu Glu Val Val Lys Ser Ser Asp Tyr Cys Glu Tyr Leu 290 295 300

Arg Pro Pro Ile Asp Ser Tyr Ser Thr Leu Asp Phe Gly Lys Phe Asn 305 310 315 320

Glu Ile Cys Glu Val Gly Tyr Gln His Gly Arg 325 330

<210> 178

<211> 289

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: UPF0028 domain sequence

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|-----|---|----------|---|---|---|
| | | | | | |

- Ile Ala Phe Gln Ser Asp Phe Ser Arg Leu Ala Arg Ile Leu Thr Gly
 1 5 10 15
- Asn Ala Ile Gly Leu Val Leu Gly Gly Gly Gly Ala Arg Gly Ala Ala 20 25 30
- His Ile Gly Val Ile Gln Ala Leu Lys Glu Val Gly Ile Pro Ile Asp 35 40 45
- Ile Val Gly Gly Thr Ser Ile Gly Ser Leu Val Gly Ala Leu Tyr Ala 50 55 60
- Cys Asp Pro Asp Ser Val Leu Val Asp Ala Arg Ala Lys Trp Phe Phe 65 70 75 80
- Ser Gly Ser Ser Ser Ile Trp Asp Arg Leu Met Asp Leu Thr Trp Pro 85 90 95
- Arg Ser Gly Leu Leu Thr Gly His Arg Phe Asn Arg Gln Val Gln Glu
 100 105 110
- Ile Phe Gly Glu Thr Leu Ile Glu Asp Cys Trp Arg Ser Phe Phe Cys
 115 120 125
- Val Ser Thr Asp Leu Ser Thr Ser Arg Gln Arg Ile His Arg Glu Gly
 130 135 140
- Pro Pro Val Cys Gln Asn Gly His Leu Leu Leu Asp Gly Gly Tyr Val 165 170 175
- Asn Asn Leu Pro Ala Asp Val Met Arg Ala Leu Gly Ala Asp Ile Val 180 185 190
- Ile Ala Val Asp Val Gly Ser Ala Asp Leu Thr Asn Leu Asp Leu Tyr 195 200 205
- Gly Phe Ser Leu Ser Gly Glu Trp Ile Leu Phe Lys Arg Trp Asn Pro 210 215 220
- Phe Gly Ala Arg Leu Arg Ile Leu Asn Met Ser Glu Ile Gln Arg Arg 225 230 235 240
- Leu Ala Tyr Val Pro Cys Val Arg Ala Leu Glu Thr Ala Lys Asn Thr 245 250 255

Val Tyr Cys Arg Tyr Leu Lys Arg Pro Ile Glu Ala Phe Asp Thr Leu 260 265 270 Asp Phe Ser Lys Phe Pro Glu Ile Pro Gln Ile Gly Val Leu Tyr Phe 280 285 275 Lys <210> 179 <211> 94 <212> PRT <213> Homo sapiens <400> 179 Ala Leu Asp Trp Val Glu Val Glu Ala Gly Arg Ala Ile Tyr Arg Gln 1 5 10 15 Gly Asp Lys Ser Asp Cys Thr Tyr Ile Met Leu Ser Gly Arg Leu Arg 25 30 20 Ser Val Ile Arg Lys Asp Asp Gly Lys Lys Arg Leu Ala Gly Glu Tyr 35 40 Gly Arg Gly Asp Leu Val Gly Val Val Glu Thr Leu Thr His Gln Ala 50 55 60 Arg Ala Thr Thr Val His Ala Val Arg Asp Ser Glu Leu Ala Lys Leu 75 80 65 70 Pro Ala Gly Ala Leu Thr Cys Ile Lys Arg Arg Tyr Pro Gln 85 90 <210> 180 <211> 94 <212> PRT <213> Artificial Sequence <220>

<223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence

<400> 180
Ala Leu Glu Glu Arg Ser Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln
1 5 10 15

Gly Asp Pro Gly Asp Ser Leu Tyr Ile Val Val Ser Gly Ser Val Glu 20 25 30

Val Tyr Arg Leu Leu Glu Asp Gly Arg Glu Gln Ile Val Gly Thr Leu 35 40 45

Gly Pro Gly Asp Leu Phe Gly Glu Leu Ala Leu Leu Thr Asn Pro Pro 50 55 60

Arg Thr Ala Thr Val Arg Ala Leu Thr Asp Cys Glu Leu Leu Arg Leu 65 70 75 80

Asp Arg Glu Asp Phe Glu Arg Leu Leu Glu Gln Tyr Pro Glu 85 90

<210> 181

<211> 89

<212> PRT

<213> Homo sapiens

<400> 181

His Val Pro Ala Gly Thr Val Val Ser Arg Gln Gly Asp Gln Asp Ala
1 5 10 15

Ser Ile Leu Phe Val Val Ser Gly Leu Leu His Val Tyr Gln Arg Lys 20 25 30

Ile Gly Ser Gln Glu Asp Thr Cys Leu Phe Leu Thr Arg Pro Gly Glu 35 40 45

Met Val Gly Gln Leu Ala Val Leu Thr Gly Glu Pro Leu Ile Phe Thr 50 55 60

Val Lys Ala Asn Arg Asp Cys Ser Phe Leu Ser Ile Ser Lys Ala His 65 70 75 80

Phe Tyr Glu Ile Met Arg Lys Gln Pro 85

<210> 182

<211> 88

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence

<400> 182

Ser Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln Gly Asp Pro Gly Asp 1 5 10 15

Ser Leu Tyr Ile Val Val Ser Gly Ser Val Glu Val Tyr Arg Leu Leu 20 25 30

Glu Asp Gly Arg Glu Gln Ile Val Gly Thr Leu Gly Pro Gly Asp Leu $35 \hspace{1cm} 40 \hspace{1cm} 45$

Phe Gly Glu Leu Ala Leu Leu Thr Asn Pro Pro Arg Thr Ala Thr Val 50 55 60

Arg Ala Leu Thr Asp Cys Glu Leu Leu Arg Leu Asp Arg Glu Asp Phe 65 70 75 80

Glu Arg Leu Leu Glu Gln Tyr Pro 85

<210> 183

<211> 101

<212> PRT

<213> Homo sapiens

<400> 183

His Ile Val Phe Val Gln Leu Gln Glu Gly Glu His Val Phe Gln Pro $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Arg Glu Pro Asp Pro Ser Ile Cys Val Val Gln Asp Gly Arg Leu Glu 20 25 30

Val Cys Ile Gln Asp Thr Asp Gly Thr Glu Val Val Val Lys Glu Val 35 40 45

Leu Ala Gly Asp Ser Val His Ser Leu Leu Ser Ile Leu Asp Ile Ile 50 55 60

Thr Gly His Ala Ala Pro Tyr Lys Thr Val Ser Val Arg Ala Ala Ile 65 70 75 80

Pro Ser Thr Ile Leu Arg Leu Pro Ala Ala Ala Phe His Gly Val Phe 85 90 95

Glu Lys Tyr Pro Glu

100

<210> 184 <211> 94 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence <400> 184 Ala Leu Glu Glu Arg Ser Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln 10 Gly Asp Pro Gly Asp Ser Leu Tyr Ile Val Val Ser Gly Ser Val Glu Val Tyr Arg Leu Leu Glu Asp Gly Arg Glu Gln Ile Val Gly Thr Leu 40 Gly Pro Gly Asp Leu Phe Gly Glu Leu Ala Leu Leu Thr Asn Pro Pro 50 55 Arg Thr Ala Thr Val Arg Ala Leu Thr Asp Cys Glu Leu Leu Arg Leu 65 70 75 80 Asp Arg Glu Asp Phe Glu Arg Leu Leu Glu Gln Tyr Pro Glu 85 90 <210> 185 <211> 115 <212> PRT <213> Homo sapiens <400> 185 Ser Phe Val Arg Gln Ile Asp Phe Ala Leu Asp Trp Val Glu Val Glu 10 Ala Gly Arg Ala Ile Tyr Arg Gln Gly Asp Lys Ser Asp Cys Thr Tyr 20 25 Ile Met Leu Ser Gly Arg Leu Arg Ser Val Ile Arg Lys Asp Asp Gly 40

Lys Lys Arg Leu Ala Gly Glu Tyr Gly Arg Gly Asp Leu Val Gly Val

50 55 60

Val Glu Thr Leu Thr His Gln Ala Arg Ala Thr Thr Val His Ala Val 65 70 75 80

Arg Asp Ser Glu Leu Ala Lys Leu Pro Ala Gly Ala Leu Thr Cys Ile 85 90 95

Lys Arg Arg Tyr Pro Gln Val Val Thr Arg Leu Ile His Leu Leu Gly
100 105 110

Glu Lys Ile 115

<210> 186

<211> 114

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence

<400> 186

Glu Glu Leu Arg Glu Leu Ala Asp Ala Leu Glu Pro Val Arg Tyr Pro 1 5 10 15

Ala Gly Glu Val Ile Ile Arg Gln Gly Asp Val Gly Asp Ser Phe Tyr 20 25 30

Ile Ile Val Ser Gly Glu Val Glu Val Tyr Lys Thr Leu Glu Asp Gly
35 40 45

Arg Glu Gln Ile Leu Gly Thr Leu Gly Pro Gly Asp Phe Phe Gly Glu 50 55 60

Leu Ala Leu Leu Thr Asn Arg Arg Arg Ala Arg Ser Ala Ala Val
65 70 75 80

Ala Leu Glu Leu Ala Lys Leu Leu Arg Ile Asp Phe Arg Asp Phe Leu 85 90 95

Gln Leu Leu Pro Glu Ile Pro Gln Leu Leu Leu Glu Leu Leu Glu
100 105 110

Leu Ala

<210> 187 <211> 123 <212> PRT <213> Homo sapiens <400> 187 Val Leu Gly His Phe Glu Lys Pro Leu Phe Leu Glu Leu Cys Lys His 5 10 Ile Val Phe Val Gln Leu Gln Glu Gly Glu His Val Phe Gln Pro Arg 20 25 Glu Pro Asp Pro Ser Ile Cys Val Val Gln Asp Gly Arg Leu Glu Val 35 Cys Ile Gln Asp Thr Asp Gly Thr Glu Val Val Lys Glu Val Leu 50 55 60 Ala Gly Asp Ser Val His Ser Leu Leu Ser Ile Leu Asp Ile Ile Thr 65 70 75 Gly His Ala Ala Pro Tyr Lys Thr Val Ser Val Arg Ala Ala Ile Pro 85 90 95 Ser Thr Ile Leu Arg Leu Pro Ala Ala Phe His Gly Val Phe Glu 100 105 110 Lys Tyr Pro Glu Thr Leu Val Arg Val Val Gln 115 120 <210> 188 <211> 118 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence <400> 188 Leu Phe Lys Ala Leu Asp Ala Glu Glu Leu Arg Glu Leu Ala Asp Ala 5 1 10 15 Leu Glu Pro Val Arg Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln Gly

25

30

Asp Val Gly Asp Ser Phe Tyr Ile Ile Val Ser Gly Glu Val Glu Val 35 40 45 .

Tyr Lys Thr Leu Glu Asp Gly Arg Glu Gln Ile Leu Gly Thr Leu Gly 50 55 60

Pro Gly Asp Phe Phe Gly Glu Leu Ala Leu Leu Thr Asn Arg Arg Arg 65 70 75 80

Ala Arg Ser Ala Ala Ala Val Ala Leu Glu Leu Ala Lys Leu Leu Arg 85 90 95

Ile Asp Phe Arg Asp Phe Leu Gln Leu Leu Pro Glu Ile Pro Gln Leu
100 105 110

Leu Leu Glu Leu Leu 115

<210> 189

<211> 89

<212> PRT

<213> Homo sapiens

<400> 189

His Val Pro Ala Gly Thr Val Val Ser Arg Gln Gly Asp Gln Asp Ala 1 5 10 15

Ser Ile Leu Phe Val Val Ser Gly Leu Leu His Val Tyr Gln Arg Lys 20 25 30

Ile Gly Ser Gln Glu Asp Thr Cys Leu Phe Leu Thr Arg Pro Gly Glu
35 40 45

Met Val Gly Gln Leu Ala Val Leu Thr Gly Glu Pro Leu Ile Phe Thr 50 55 60

Val Lys Ala Asn Arg Asp Cys Ser Phe Leu Ser Ile Ser Lys Ala His 65 70 75 80

Phe Tyr Glu Ile Met Arg Lys Gln Pro 85

<210> 190

<211> 90

<212> PRT

<213> Artificial Sequence <220> <400> 190 20 40

<223> Description of Artificial Sequence: Cyclic nucleotide-binding domain sequence

Arg Tyr Pro Ala Gly Glu Val Ile Ile Arg Gln Gly Asp Val Gly Asp 10

Ser Phe Tyr Ile Ile Val Ser Gly Glu Val Glu Val Tyr Lys Thr Leu 25

Glu Asp Gly Arg Glu Gln Ile Leu Gly Thr Leu Gly Pro Gly Asp Phe

Phe Gly Glu Leu Ala Leu Leu Thr Asn Arg Arg Arg Ala Arg Ser Ala

Ala Ala Val Ala Leu Glu Leu Ala Lys Leu Leu Arg Ile Asp Phe Arg 70 75

Asp Phe Leu Gln Leu Leu Pro Glu Ile Pro 85

<210> 191

<211> 330

<212> PRT

<213> Homo sapiens

<400> 191

Met Arg Arg Pro Ser Val Arg Ala Ala Gly Leu Val Leu Cys Thr Leu 10 1 15

Cys Tyr Leu Leu Val Gly Ala Ala Val Phe Asp Ala Leu Glu Ser Glu 20 25

Ala Glu Ser Gly Arg Gln Arg Leu Leu Val Gln Lys Arg Gly Ala Leu 40

Arg Arg Lys Phe Gly Phe Ser Ala Glu Asp Tyr Arg Glu Leu Glu Arg 55

Leu Ala Leu Gln Ala Glu Pro His Arg Ala Gly Arg Gln Trp Lys Phe 70 75

Pro Gly Ser Phe Tyr Phe Ala Ile Thr Val Ile Thr Thr Ile Glu Tyr

| Gly | His | Ala | Ala 100 | Pro | Gly | Thr | Asp | Ser 105 | Gly | Lys | Val | Phe | Cys 110 | Met | Phe |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Tyr | Ala | Leu 115 | Leu | Gly | Ile | Pro | Leu 120 | Thr | Leu | Val | Thr | Phe 125 | Gln | Ser | Leu |
| Gly | Glu 130 | Arg | Leu | Asn | Ala | Val 135 | Val | Arg | Arg | Leu | Leu 140 | Leu | Ala | Ala | Lys |
| Cys 145 | Cys | Leu | Gly | Leu | Arg 150 | Trp | Thr | Cys | Val | Ser 155 | Thr | Glu | Asn | Leu | Val 160 |
| Val | Ala | Gly | Leu | Leu 165 | Ala | Cys | Ala | Ala | Thr 170 | Leu | Ala | Leu | Gly | Ala 175 | Val |
| Ala | Phe | Ser | His 180 | Phe | Glu | Gly | Trp | Thr 185 | Phe | Phe | His | Ala | Tyr 190 | Tyr | Tyr |
| Cys | Phe | Ile 195 | Thr | Leu | Thr | Thr | Ile 200 | Gly | Phe | Gly | Asp | Phe 205 | Val | Ala | Leu |
| Gln | Ser 210 | Gly | Glu | Ala | Leu | Gln 215 | Arg | Lys | Leu | Pro | Tyr 220 | Val | Ala | Phe | Ser |
| Phe 225 | Leu | Tyr | Ile | Leu | Leu 230 | Gly | Leu | Thr | Val | Ile 235 | Gly | Ala | Phe | Leu | Asn 240 |
| Leu | Val | Val | Leu | Arg 245 | Phe | Leu | Val | Ala | Ser 250 | Ala | Asp | Trp | Pro | Glu 255 | Arg |
| Ala | Ala | Arg | Thr 260 | Pro | Ser | Pro | Arg | Pro 265 | Pro | Gly | Ala | Pro | Glu 270 | Ser | Arg |
| Gly | Leu | Trp 275 | Leu | Pro | Arg | Arg | Pro 280 | Ala | Arg | Ser | Val | Gly 285 | Ser | Ala | Ser |
| Val | Phe 290 | Cys | His | Val | His | Lys 295 | Leu | Glu | Arg | Cys | Ala 300 | Arg | Asp | Asn | Leu |
| Gly 305 | Phe | Ser | Pro | Pro | Ser 310 | Ser | Pro | Gly | Val | Val 315 | Arg | Gly | Gly | Gln | Ala 320 |
| Pro | Arg | Leu | Gly | Ala 325 | Arg | Trp | Lys | Ser | Ile 330 | | | | | | |
| | | | | | | | | | | | | | | | |

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<210> 192
<211> 330
<212> PRT
<213> Homo sapiens
<400> 192
Met Arg Arg Pro Ser Val Arg Ala Ala Gly Leu Val Leu Cys Thr Leu
                  5
                                      10
                                                          15
Cys Tyr Leu Leu Val Gly Ala Ala Val Phe Asp Ala Leu Glu Ser Glu
             20
                                 25
Ala Glu Ser Gly Arg Gln Arg Leu Leu Val Gln Lys Arg Gly Ala Leu
                                                  45
         35
                             40
Arg Arg Lys Phe Gly Phe Ser Ala Glu Asp Tyr Arg Glu Leu Glu Arg
                         55
Leu Ala Leu Gln Ala Glu Pro His Arg Ala Gly Arg Gln Trp Lys Phe
                     70
                                          75
 65
Pro Gly Ser Phe Tyr Phe Ala Ile Thr Val Ile Thr Thr Ile Gly Tyr
                 85
                                      90
Gly His Ala Ala Pro Gly Thr Asp Ser Gly Lys Val Phe Cys Met Phe
            100
                                105
Tyr Ala Leu Leu Gly Ile Pro Leu Thr Leu Val Thr Phe Gln Ser Leu
        115
                            120
                                                 125
Gly Glu Arg Leu Asn Ala Val Val Arg Arg Leu Leu Ala Ala Lys
                        135
Cys Cys Leu Gly Leu Arg Trp Thr Cys Val Ser Thr Glu Asn Leu Val
                    150
                                        155
Val Ala Gly Leu Leu Ala Cys Ala Ala Thr Leu Ala Leu Gly Ala Val
                165
                                    170
Ala Phe Ser His Phe Glu Gly Trp Thr Phe Phe His Ala Tyr Tyr Tyr
            180
                                185
```

Gln Ser Gly Glu Ala Leu Gln Arg Lys Leu Pro Tyr Val Ala Phe Ser 210 215 220

Cys Phe Ile Thr Leu Thr Thr Ile Gly Phe Gly Asp Phe Val Ala Leu

205

200

| 225 | Tyr | Ile | Leu | Leu 230 | Gly | Leu | Thr | Val | Ile 235 | Gly | Ala | Phe | Leu | Asn 240 |
|---|---|--------------------|-------------------------------|--------------------------------|-------------------------|--------------------------------|--------------------------------|-------------------------|--------------------------------|--------------------------------|-------------------------|--------------------------------|--------------------------------|--------------------------------|
| Leu Val | Val | Leu | Arg 245 | Phe | Leu | Val | Ala | Ser 250 | Ala | Asp | Trp | Pro | Glu 255 | Arg |
| Ala Ala | Arg | Pro 260 | Pro | Ser | Pro | Arg | Pro 265 | Pro | Gly | Ala | Pro | Glu 270 | Ser | Arg |
| Gly Leu | Trp 275 | Leu | Pro | Arg | Arg | Pro 280 | Ala | Arg | Ser | Val | Gly 285 | Ser | Ala | Ser |
| Val Phe 290 | Cys | His | Val | His | Lys 295 | Leu | Glu | Arg | Cys | Ala 300 | Arg | Asp | Asn | Leu |
| Gly Phe 305 | Ser | Pro | Pro | Ser 310 | Ser | Pro | Gly | Val | Val 315 | Arg | Gly | Gly | Gln | Ala 320 |
| Pro Arg | Pro | Gly | Ala 325 | Arg | Trp | Lys | Ser | Ile 330 | | | | | | |
| <210> 1 <211> 3 | | | | | | | | | | | | | | |
| 4010\ D | ייים | | | | | | | | | | | | | |
| <212> P <213> H | | sapie | ens | | | | | | | | | | | |
| | omo : | sapie | ens | | | | | | | | | | | |
| <213> H | omo : 93 | _ | | Val | Arg | Ala | Ala | Gly 10 | Leu | Val | Leu | Cys | Thr 15 | Leu |
| <213> H <400> 1 Met Arg | omo s 93 Arg | Pro | Ser 5 | | | | | 10 | | | | | 15 | |
| <213> H <400> 1 Met Arg 1 | omo s 93 Arg Leu | Pro | Ser 5 Val | Gly | Ala | Ala | Val 25 | 10 Phe | Asp | Ala | Leu | Glu 30 | 15 Ser | Glu |
| <213> H <400> 1 Met Arg 1 Cys Tyr | omo s 93 Arg Leu Ser 35 | Pro Leu 20 Gly | Ser 5 Val Arg | Gly | Ala | Ala Leu 40 | Val 25 Leu | 10 Phe Val | Asp | Ala Lys | Leu Arg 45 | Glu 30 Gly | 15 Ser Ala | Glu Leu |
| <213> H <400> 1 Met Arg 1 Cys Tyr Ala Glu Arg Arg | omo s 93 Arg Leu Ser 35 Lys | Pro Leu 20 Gly Phe | Ser 5 Val Arg | Gly Gln Phe | Ala Arg Ser 55 | Ala Leu 40 Ala | Val 25 Leu Glu | 10 Phe Val Asp | Asp Gln Tyr | Ala Lys Arg 60 | Leu Arg 45 Glu | Glu 30 Gly Leu | 15 Ser Ala Glu | Glu Leu Arg |
| <213> H <400> 1 Met Arg | omo s 93 Arg Leu Ser 35 Lys Leu | Pro Leu 20 Gly Phe | Ser 5 Val Arg Gly | Gly Gln Phe Glu 70 | Ala Arg Ser 55 | Ala Leu 40 Ala His | Val 25 Leu Glu Arg | 10 Phe Val Asp | Asp Gln Tyr Gly 75 | Ala Lys Arg 60 Arg | Leu Arg 45 Glu | Glu 30 Gly Leu Trp | 15 Ser Ala Glu Lys | Glu Leu Arg Phe 80 |

Tyr Ala Leu Leu Gly Ile Pro Leu Thr Leu Val Thr Phe Gln Ser Leu
115 120 125

Gly Glu Arg Leu Asn Ala Val Val Arg Arg Leu Leu Ala Ala Lys 130 135 140

Val Ala Gly Leu Leu Ala Cys Ala Ala Thr Leu Ala Leu Gly Ala Val 165 170 175

Ala Phe Ser His Phe Glu Gly Trp Thr Phe Phe His Ala Tyr Tyr Tyr 180 185 190

Cys Phe Ile Thr Leu Thr Thr Ile Gly Phe Gly Asp Phe Val Ala Leu 195 200 205

Gln Ser Gly Glu Ala Leu Gln Arg Lys Leu Pro Tyr Val Ala Phe Ser 210 215 220

Phe Leu Tyr Ile Leu Leu Gly Leu Thr Val Ile Gly Ala Phe Leu Asn 225 230 235 240

Leu Val Val Leu Arg Phe Leu Val Ala Ser Ala Asp Trp Pro Glu Arg
245 250 255

Ala Ala Arg Thr Pro Ser Pro Arg Pro Pro Gly Ala Pro Glu Ser Arg
260 265 270

Gly Leu Trp Leu Pro Arg Arg Pro Ala Arg Ser Val Gly Ser Ala Ser 275 280 285

Val Phe Cys His Val His Lys Leu Glu Arg Cys Ala Arg Asp Asn Leu 290 295 300

Gly Phe Ser Pro Pro Ser Ser Pro Gly Val Val Arg Gly Gln Ala 305 310 315 320

Pro Arg Leu Gly Ala Arg Trp Lys Ser Ile 325 330

<210> 194

<211> 374

<212> PRT

<213> Homo sapiens

| <400 |)> 19 | 94 | | | | | | | | | | | | | |
|-----------|------------|------------|------------|-----------|-----------|------------|------------|------------|-----------|-----------|------------|------------|------------|-----------|-----------|
| Met 1 | Lys | Arg | Gln | Asn 5 | Val | Arg | Thr | Leu | Ser 10 | Leu | Ile | Val | Cys | Thr 15 | Phe |
| Thr | Tyr | Leu | Leu 20 | Val | Gly | Ala | Ala | Val 25 | Phe | Asp | Ala | Leu | Glu 30 | Ser | Asp |
| His | Glu | Met 35 | Arg | Glu | Glu | Glu | Lys 40 | Leu | Lys | Ala | Glu | Glu 45 | Ile | Arg | Ile |
| Lys | Gly 50 | Lys | Tyr | Asn | Ile | Ser 55 | Ser | Glu | Asp | Tyr | Arg 60 | Gln | Leu | Glu | Leu |
| Val 65 | Ile | Leu | Gln | Ser | Glu 70 | Pro | His | Arg | Ala | Gly 75 | Val | Gln | Trp | Lys | Phe 80 |
| Ala | Gly | Ser | Phe | Tyr 85 | Phe | Ala | Ile | Thr | Val 90 | Ile | Thr | Thr | Ile | Gly 95 | Tyr |
| Gly | His | Ala | Ala 100 | Pro | Gly | Thr | Asp | Ala 105 | Gly | Lys | Ala | Phe | Cys 110 | Met | Phe |
| Tyr | Ala | Val 115 | Leu | Gly | Ile | Pro | Leu 120 | Thr | Leu | Val | Met | Phe 125 | Gln | Ser | Leu |
| Gly | Glu 130 | Arg | Met | Asn | Thr | Phe 135 | Val | Arg | Tyr | Leu | Leu 140 | Lys | Arg | Ile | Lys |

Lys Cys Cys Gly Met Arg Asn Thr Asp Val Ser Met Glu Asn Met Val 145 150 155 160

Thr Val Gly Phe Phe Ser Cys Met Gly Thr Leu Cys Ile Gly Ala Ala 165 170 175

Ala Phe Ser Gln Cys Glu Glu Trp Ser Phe Phe His Ala Tyr Tyr Tyr 180 185 190

Cys Phe Ile Thr Leu Thr Thr Ile Gly Phe Gly Asp Tyr Val Ala Leu 195 200 205

Gln Thr Lys Gly Ala Leu Gln Lys Lys Pro Leu Tyr Val Ala Phe Ser 210 215 220

Phe Met Tyr Ile Leu Val Gly Leu Thr Val Ile Gly Ala Phe Leu Asn 225 230 235 240

Leu Val Val Leu Arg Phe Leu Thr Met Asn Ser Glu Asp Glu Arg Arg

Asp Ala Glu Glu Arg Ala Ser Leu Ala Gly Asn Arg Asn Ser Met Val 260 265 270

Ile His Ile Pro Glu Glu Pro Arg Pro Ser Arg Pro Arg Tyr Lys Ala 275 280 285

Asp Val Pro Asp Leu Gln Ser Val Cys Ser Cys Thr Cys Tyr Arg Ser 290 295 300

Gln Asp Tyr Gly Gly Arg Ser Val Ala Pro Gln Asn Ser Phe Ser Ala 305 310 315 320

Lys Leu Ala Pro His Tyr Phe His Ser Ile Ser Tyr Lys Ile Glu Glu 325 330 335

Ile Ser Pro Ser Thr Leu Lys Asn Ser Leu Phe Pro Ser Pro Ile Ser 340 345 350

Ser Ile Ser Pro Gly Leu His Ser Phe Thr Asp His Gln Arg Leu Met 355 360 365

Lys Arg Arg Lys Ser Val 370

<210> 195

<211> 387

<212> PRT

<213> Cavia porcellus

<400> 195

Met Lys Lys Gln Asn Val Arg Thr Leu Ser Leu Ile Ala Cys Thr Phe 1 5 10 15

Thr Tyr Leu Leu Val Gly Ala Ala Val Phe Asp Ala Leu Glu Ser Asp 20 25 30

His Glu Met Arg Glu Glu Glu Lys Leu Lys Ala Glu Glu Ile Arg Ile 35 40 45

Arg Gly Lys Tyr Asn Ile Ser Thr Glu Asp Tyr Arg Gln Leu Glu Leu 50 55 60

Val Ile Leu Gln Ser Glu Pro His Arg Ala Gly Val Gln Trp Lys Phe 65 70 75 80

| Ala | Gly | Ser | Phe | Tyr 85 | Phe | Ala | Ile | Thr | Val 90 | Ile | Thr | Thr | Ile | Gly 95 | Tyr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly | His | Ala | Ala 100 | Pro | Gly | Thr | Asp | Ala 105 | Gly | Lys | Ala | Phe | Cys 110 | Met | Phe |
| Tyr | Ala | Val 115 | Leu | Gly | Ile | Pro | Leu 120 | Thr | Leu | Val | Met | Phe 125 | Gln | Ser | Leu |
| Gly | Glu 130 | Arg | Met | Asn | Thr | Phe 135 | Val | Arg | Tyr | Leu | Leu 140 | Lys | Arg | Ile | Lys |
| Lys 145 | Cys | Cys | Gly | Met | Arg 150 | Asn | Thr | Glu | Val | Ser 155 | Met | Glu | Asn | Met | Val 160 |
| Thr | Val | Gly | Phe | Phe 165 | Ser | Cys | Met | Gly | Thr 170 | Leu | Cys | Ile | Gly | Ala 175 | Ala |
| Ala | Phe | Ser | Gln 180 | Cys | Glu | Glu | Trp | Ser 185 | Phe | Phe | His | Ala | Tyr 190 | Tyr | Tyr |
| Cys | Phe | Ile 195 | Thr | Leu | Thr | Thr | Ile 200 | | Phe | Gly | Asp | Tyr 205 | Val | Ala | Leu |
| Gln | Ser 210 | Lys | Gly | Ala | Leu | Gln 215 | Arg | Lys | Pro | Phe | Tyr 220 | Val | Ala | Phe | Ser |
| Phe 225 | Met | Tyr | Ile | Leu | Val 230 | Gly | Leu | Thr | Val | Ile 235 | Gly | Ala | Phe | Leu | Asn 240 |
| Leu | Val | Val | Leu | Arg 245 | Phe | Leu | Thr | Met | Asn 250 | Ser | Asp | Glu | Glu | Arg 255 | Gly |
| Glu | Gly | Glu | Glu 260 | Gly | Ala | Ala | Leu | Pro 265 | Gly | Asn | Pro | Ser | Ser 270 | Val | Val |
| Thr | His | Ile 275 | Ser | Glu | Glu | Ala | Arg 280 | Gln | Val | Arg | Gln | Arg 285 | Tyr | Arg | Gly |
| Glu | Gly 290 | Gly | Asp | Leu | Gln | Ser 295 | Val | Суѕ | Ser | Cys | Ala 300 | Cys | Tyr | Arg | Ser |
| Gln 305 | Pro | Gln | Asn | Phe | Gly 310 | His | Lys | Leu | Glu | Arg 315 | Cys | Ala | Arg | Asp | Asn 320 |
| Leu | Gly | Phe | Ser | Pro 325 | Pro | Ser | Ser | Pro | Gly 330 | Val | Val | Ala | Thr | Leu 335 | Ala |

Pro Gln Pro Leu His Ser Ile Ser Cys Arg Ile Glu Glu Ile Ser Pro 340 345 350

Ser Thr Leu Lys Asn Ser Leu Phe Pro Ser Pro Ile Ser Ser Val Ser 355 360 365

Pro Gly Leu His Ser Phe Gly Asp Asn His Arg Leu Met Leu Arg Arg 370 375 380

Lys Ser Val 385

<210> 196

<211> 257

<212> PRT

<213> Xenopus laevis

<400> 196

Met Gly Arg Val Ile Arg Gly Gln Arg Lys Gly Ala Gly Ser Val Phe
1 5 10 15

Lys Ala His Val Lys His Arg Lys Gly Ala Ala Lys Leu Arg Ala Ile 20 25 30

Asp Phe Ala Glu Arg Asn Gly Tyr Ile Lys Gly Ile Val Lys Asp Ile 35 40 45

Ile His Asp Pro Gly Arg Gly Ala Pro Leu Ala Lys Val Ala Phe Arg 50 55 60

Asp Pro Tyr Arg Phe Lys Lys Arg Thr Glu Leu Phe Val Ala Ala Glu 65 70 75 80

Gly Ile His Thr Gly Gln Phe Val Tyr Cys Gly Lys Lys Ala Gln Leu 85 90 95

Asn Ile Gly Asn Val Leu Pro Val Gly Thr Met Pro Glu Gly Thr Ile 100 · 105 110

Val Cys Cys Val Glu Glu Lys Pro Gly Asp Arg Gly Lys Leu Ala Arg 115 120 125

Ala Ser Gly Asn Tyr Ala Thr Val Ile Ser His Asn Pro Glu Thr Lys 130 135 140

Lys Thr Arg Val Lys Leu Pro Ser Gly Ser Lys Lys Val Ile Ser Ser 145 150 155 160

Ala Asn Arg Ala Ile Val Gly Val Val Ala Gly Gly Gly Arg Ile Asp 165 170 175

Lys Pro Ile Leu Lys Ala Gly Arg Ala Tyr His Lys Tyr Lys Ala Lys 180 185 190

Arg Asn Cys Trp Pro Arg Val Arg Gly Val Ala Met Asn Pro Val Glu
195 200 205

His Pro Phe Gly Gly Gly Asn His Gln His Ile Gly Lys Pro Ser Thr 210 215 220

Ile Arg Arg Asp Ala Pro Ala Gly Arg Lys Val Gly Leu Ile Ala Ala 225 230 235 240

Arg Arg Thr Gly Arg Leu Arg Gly Thr Lys Thr Val Gln Glu Lys Glu 245 250 255

Asn

<210> 197

<211> 257

<212> PRT

<213> Homo sapiens

<400> 197

Met Gly Arg Val Ile Arg Gly Gln Arg Lys Gly Ala Gly Ser Val Phe
1 5 10 15

Arg Ala His Val Lys His Arg Lys Gly Ala Ala Arg Leu Arg Ala Val 20 25 30

Asp Phe Ala Glu Arg His Gly Tyr Ile Lys Gly Ile Val Lys Asp Ile 35 40 45

Ile His Asp Pro Gly Arg Gly Ala Pro Leu Ala Lys Val Val Phe Arg 50 55 60

Asp Pro Tyr Arg Phe Lys Lys Arg Thr Glu Leu Phe Ile Ala Ala Glu 65 70 75 80

Gly Ile His Thr Gly Gln Phe Val Tyr Cys Gly Lys Lys Ala Gln Leu 85 90 95

Asn Ile Gly Asn Val Leu Pro Val Gly Thr Met Pro Glu Gly Thr Ile

100 105 110

Val Cys Cys Leu Glu Glu Lys Pro Gly Asp Arg Gly Lys Leu Ala Arg 115 120 125

Ala Ser Gly Asn Tyr Ala Thr Val Ile Ser His Asn Pro Glu Thr Lys 130 135 140

Lys Thr Arg Val Lys Leu Pro Ser Gly Ser Lys Lys Val Ile Ser Ser 145 150 155 160

Ala Asn Arg Ala Val Val Gly Val Val Ala Gly Gly Gly Arg Ile Asp 165 170 175

Lys Pro Ile Leu Lys Ala Gly Arg Ala Tyr His Lys Tyr Lys Ala Lys 180 185 190

Arg Asn Cys Trp Pro Arg Val Arg Gly Val Ala Met Asn Pro Val Glu
195 200 205

His Pro Phe Gly Gly Gly Asn His Gln His Ile Gly Lys Pro Ser Thr 210 215 220

Ile Arg Arg Asp Ala Pro Ala Gly Arg Lys Val Gly Leu Ile Ala Ala 225 230 235 240

Arg Arg Thr Gly Arg Leu Arg Gly Thr Lys Thr Val Gln Glu Lys Glu 245 250 255

Asn

<210> 198

<211> 257

<212> PRT

<213> Homo sapiens

<400> 198

Met Gly Arg Val Ile Arg Gly Gln Arg Lys Gly Ala Gly Ser Val Phe 1 5 10 15

Arg Ala His Val Lys His Arg Lys Gly Ala Ala Arg Leu Arg Ala Val 20 25 30

Asp Phe Ala Glu Arg His Gly Tyr Ile Lys Gly Ile Val Lys Asp Ile 35 40 45

Ile His Asp Pro Gly Arg Gly Ala Pro Leu Ala Lys Val Val Phe Arg
50 55 60

Asp Pro Tyr Arg Phe Lys Lys Arg Thr Glu Leu Phe Ile Ala Ala Glu 65 70 75 80

Gly Ile His Thr Gly Gln Phe Val Tyr Cys Gly Lys Lys Ala Gln Leu 85 90 95

Asn Val Gly Asn Val Leu Pro Val Gly Thr Met Pro Glu Gly Thr Ile 100 105 110

Val Cys Cys Leu Glu Glu Lys Pro Gly Asp Arg Gly Lys Leu Ala Arg 115 120 125

Ala Ser Gly Asn Tyr Ala Thr Val Ile Ser His Asn Pro Glu Thr Lys 130 135 140

Lys Thr Arg Val Lys Leu Pro Ser Gly Ser Lys Lys Val Ile Ser Ser 145 150 155 160

Ala Asn Arg Ala Val Val Gly Val Val Ala Gly Gly Gly Arg Ile Asp 165 170 175

Lys Pro Ile Leu Lys Ala Gly Arg Ala Tyr His Lys Tyr Lys Ala Lys
180 185 190

Arg Asn Cys Trp Pro Arg Val Arg Gly Val Ala Met Asn Pro Val Glu
195 200 205

His Pro Phe Gly Gly Gly Asn His Gln His Ile Gly Lys Pro Ser Thr 210 215 220

Ile Arg Arg Asp Ala Pro Ala Gly Arg Lys Val Gly Leu Ile Ala Ala 225 230 235 240

Arg Arg Thr Gly Arg Leu Arg Gly Thr Lys Thr Val Gln Glu Lys Glu 245 250 255

Asn

<210> 199

<211> 257

<212> PRT

<213> Ictalurus punctatus

| <4 | <00 | 199 |
|----|-----|-----|
| | | |

- Met Gly Arg Val Ile Arg Ala Gln Arg Lys Gly Ala Gly Ser Val Phe 1 5 10 15
- Lys Ala His Val Lys His Arg Lys Gly Ala Ala Lys Leu Arg His Ile 20 25 30
- Asp Phe Ala Glu Arg His Gly Tyr Ile Lys Gly Ile Val Lys Asp Ile 35 40 45
- Ile His Asp Pro Gly Arg Gly Thr Pro Leu Ala Lys Val Val Phe Arg
 50 55 60
- Asp Pro Tyr Arg Phe Lys Lys Arg Thr Glu Leu Phe Ile Ala Ala Glu 65 70 75 80
- Gly Ile His Thr Gly Gln Phe Val Phe Cys Gly Lys Lys Ala Gln Leu 85 90 95
- Asn Ile Gly Asn Val Leu Pro Val Gly Val Met Pro Glu Gly Thr Ile 100 105 110
- Ile Cys Cys Leu Glu Glu Lys Pro Gly Asp Arg Gly Lys Leu Ala Arg 115 120 125
- Ala Ser Gly Asn Tyr Ala Thr Val Ile Ser His Asn Pro Glu Thr Lys 130 135 140
- Lys Ser Arg Val Lys Leu Pro Ser Gly Ala Lys Lys Val Ile Ser Ser 145 150 155 160
- Thr Asn Arg Ala Val Val Gly Val Val Ala Gly Gly Gly Arg Ile Asp 165 170 175
- Lys Pro Ile Leu Lys Ala Gly Arg Ala Tyr His Lys Tyr Lys Val Lys 180 185 190
- Arg Asn Cys Trp Pro Arg Val Arg Gly Val Ala Met Asn Pro Val Glu 195 200 205
- His Pro Phe Gly Gly Gly Asn His Gln His Ile Gly Lys Pro Ser Thr 210 215 220
- Ile Arg Arg Asp Val Pro Ala Gly Arg Lys Val Gly Leu Ile Ala Ala 225 230 235 240
- Arg Arg Thr Gly Arg Leu Arg Gly Thr Lys Thr Val Gln Glu Lys Glu 245 250 255

| <211 <212 |)> 20 L> 21 2> PF 3> Ho | . 4 RT | sapie | ens | | | | | | | | | | | |
|--------------|----------------------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| |)> 2(Glu | | Asp | Ile 5 | Ile | His | Asp | Pro | Gly 10 | Arg | Gly | Ala | Pro | Leu 15 | Ala |
| Lys | Val | Val | Phe 20 | Arg | Asp | Pro | Tyr | Arg 25 | Phe | Lys | Lys | Arg | Thr 30 | Glu | Leu |
| Phe | Ile | Ala 35 | Ala | Glu | Gly | Ile | His 40 | Thr | Gly | Gln | Phe | Val 45 | Tyr | Cys | Gly |
| Lys | Lys 50 | Ala | Gln | Leu | Asn | Ile 55 | Gly | Asn | Val | Leu | Pro 60 | Val | Gly | Thr | Met |
| Pro 65 | Glu | Gly | Thr | Ile | Val 70 | Cys | Cys | Leu | Glu | Glu 75 | Lys | Pro | Gly | Asp | Arg 80 |
| Gly | Lys | Leu | Ala | Arg 85 | Ala | Ser | Gly | Asn | Tyr 90 | Ala | Thr | Val | Ile | Ser 95 | His |
| Asn | Pro | Glu | Thr 100 | Lys | Lys | Thr | Arg | Val 105 | Lys | Leu | Pro | Ser | Gly 110 | Ser | Lys |
| Lys | Val | Ile 115 | Ser | Ser | Ala | Asn | Arg 120 | Ala | Val | Val | Gly | Val 125 | Val | Ala | Gly |
| Gly | Gly 130 | Arg | Ile | Asp | Lys | Pro 135 | Ile | Leu | Lys | Ala | Gly 140 | Arg | Ala | Tyr | His |
| Lys 145 | Tyr | Lys | Ala | Lys | Arg 150 | Asn | Cys | Trp | Pro | Arg 155 | Val | Arg | Gly | Val | Ala 160 |
| Met | Asn | Pro | Val | Glu 165 | His | Pro | Phe | Gly | Gly 170 | Gly | Asn | His | Gln | His 175 | Ile |

Gly Lys Pro Ser Thr Ile Arg Arg Asp Ala Pro Ala Gly Arg Lys Val

Gly Leu Ile Ala Ala Arg Arg Thr Gly Arg Leu Arg Gly Thr Lys Thr

185

195 200 205

Val Gln Glu Lys Glu Asn 210

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<211> 190

<212> PRT

<213> Homo sapiens

<400> 201

Gly Ser Val Phe Arg Ala His Val Lys His Arg Lys Gly Ala Ala Arg 1 5 10 15

Leu Arg Ala Val Asp Phe Ala Glu Arg His Gly Tyr Ile Lys Gly Ile 20 25 30

Val Lys Ala Gln Leu Asn Ile Gly Asn Val Leu Pro Val Gly Thr Met 35 40 45

Pro Glu Gly Thr Ile Val Cys Cys Leu Glu Glu Lys Pro Gly Asp Arg 50 55 60

Gly Lys Leu Ala Arg Ala Ser Gly Asn Tyr Ala Thr Val Ile Ser His 65 70 75 80

Asn Pro Glu Thr Lys Lys Thr Arg Val Lys Leu Pro Ser Gly Ser Lys 85 90 95

Lys Val Ile Ser Ser Ala Asn Arg Ala Val Val Gly Val Val Ala Gly
100 105 110

Gly Gly Arg Ile Asp Lys Pro Ile Leu Lys Ala Gly Arg Ala Tyr His
115 120 125

Lys Tyr Lys Ala Lys Arg Asn Cys Trp Pro Arg Val Arg Gly Val Ala 130 135 140

Met Asn Pro Val Glu His Pro Phe Gly Gly Asn His Gln His Ile 145 150 155 160

Gly Lys Pro Ser Thr Ile Arg Arg Asp Ala Pro Ala Gly Arg Lys Val 165 170 175

Gly Leu Ile Ala Ala Arg Arg Thr Gly Arg Leu Arg Gly Thr 180 185 190 <210> 202

<211> 229

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ribosomal Proteins L2 domain sequence

<400> 202

Gly Arg Asn Asn Arg Gly His Ile Thr Arg Arg His Arg Gly Gly 1 5 10 15

His Lys Arg Leu Tyr Arg Ala Ile Asp Phe Lys Arg Arg Lys Gly Tyr
20 25 30

Ile Lys Gly Thr Val Lys Arg Ile Glu Tyr Asp Pro Asn Arg Ser Ala 35 40 45

Pro Ile Ala Leu Val Val Tyr Ser Asp Pro Gly Glu Lys Arg Tyr Ile 50 55 60

Leu Ala Pro Glu Gly Leu His Val Gly Asp Thr Ile Tyr Ser Gly Lys
65 70 75 80

Asn Ala Thr Ile Lys Ile Gly Asn Val Leu Pro Leu Gly Glu Ile Pro 85 90 95

Glu Gly Thr Ile Ile His Asn Val Glu Glu Lys Pro Gly Asp Gly Gly 100 105 110

Gln Leu Ala Arg Ala Ala Gly Thr Tyr Ala Gln Ile Leu Ala His Asp 115 120 125

Gly Asp Lys Lys Thr Arg Val Lys Leu Pro Ser Gly Glu Lys Arg Arg 130 135 140

Val Ser Ser Glu Cys Arg Ala Thr Ile Gly Val Val Ala Asn Gly Gly 145 150 155 160

Arg Ile Asp Lys Pro Leu Gly Lys Ala Gly Arg Ala Arg Trp Leu Gly 165 170 175

Lys Arg Pro Arg Val Arg Gly Val Ala Met Asn Pro Val Asp His Pro 180 185 190

His Gly Gly Glu Gly Arg His Pro Ile Gly Arg Lys Ser Pro Val

195 200 205

Thr Pro Trp Gly Lys Lys Ala Leu Gly Ile Ala Thr Arg Arg Thr Lys 210 215 220

Arg Leu Ser Asp Lys 225

<210> 203

<211> 519

<212> PRT

<213> Homo sapiens

<400> 203

Met Ser Val Ser Val Leu Ser Pro Ser Arg Leu Leu Gly Asp Val Ser 1 5 10 15

Gly Ile Leu Gln Ala Ala Ser Leu Leu Ile Leu Leu Leu Leu Leu Ile 20 25 30

Lys Ala Val Gln Leu Tyr Leu His Arg Gln Trp Leu Leu Lys Ala Leu 35 40 45

Gln Gln Phe Pro Cys Pro Pro Ser His Trp Leu Phe Gly His Ile Gln 50 55 60

Glu Leu Gln Gln Asp Gln Glu Leu Gln Arg Ile Gln Lys Trp Val Glu 65 70 75 80

Thr Phe Pro Ser Ala Cys Pro His Trp Leu Trp Gly Gly Lys Val Arg 85 90 95

Val Gln Leu Tyr Asp Pro Asp Tyr Met Lys Val Ile Leu Gly Arg Ser 100 105 110

Asp Pro Lys Ser His Gly Ser Tyr Arg Phe Leu Ala Pro Trp Ile Gly
115 120 125

Tyr Gly Leu Leu Leu Asn Gly Gln Thr Trp Phe Gln His Arg Arg 130 135 140

Leu Met Ala Asp Ser Val Arg Val Met Leu Asp Lys Trp Glu Glu Leu 165 170 175

| Leu | Gly | Gln | Asp 180 | Ser | Pro | Leu | Glu | Val 185 | Phe | Gln | His | Val | Ser 190 | Leu | Met |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Thr | Leu | Asp 195 | Thr | Ile | Met | Lys | Cys 200 | Ala | Phe | Ser | His | Gln 205 | Gly | Ser | Ile |
| Gln | Val 210 | Asp | Arg | Asn | Ser | Gln 215 | Ser | Tyr | Ile | Gln | Ala 220 | Ile | Ser | Asp | Leu |
| Asn 225 | Asn | Leu | Val | Phe | Ser 230 | Arg | Val | Arg | Asn | Ala 235 | Phe | His | Gln | Asn | Asp 240 |
| Thr | Ile | Tyr | Ser | Leu 245 | Thr | Ser | Ala | Gly | Arg 250 | Trp | Thr | His | Arg | Ala 255 | Cys |
| Gln | Leu | Ala | His 260 | Gln | His | Thr | Asp | Gln 265 | Val | Ile | Gln | Leu | Arg 270 | Lys | Ala |
| Gln | Leu | Gln 275 | Lys | Glu | Gly | Glu | Leu 280 | Glu | Lys | Ile | Lys | Arg 285 | Lys | Arg | His |
| Leu | Asp 290 | Phe | Leu | Asp | Ile | Leu 295 | Leu | Leu | Ala | Lys | Met 300 | Glu | Asn | Gly | Ser |
| Ile 305 | Leu | Ser | Asp | Lys | Asp 310 | Leu | Arg | Ala | Glu | Val 315 | Asp | Thr | Phe | Met | Phe 320 |
| Glu | Gly | His | Asp | Thr 325 | Thr | Ala | Ser | Gly | Ile 330 | Ser | Trp | Ile | Leu | Tyr 335 | Ala |
| Leu | Ala | Thr | His 340 | Pro | Lys | His | Gln | Glu 345 | Arg | Cys | Arg | Glu | Glu 350 | Ile | His |
| Ser | Leu | Leu 355 | Gly | Asp | Gly | Ala | Ser 360 | Ile | Thr | Trp | Asn | His 365 | Leu | Asp | Gln |
| Met | Pro 370 | Tyr | Thr | Thr | Met | Cys 375 | Ile | Lys | Glu | Ala | Leu 380 | Arg | Leu | Tyr | Pro |
| Pro 385 | Val | Pro | Gly | Ile | Gly 390 | Arg | Glu | Leu | Ser | Thr 395 | Pro | Val | Thr | Phe | Pro 400 |
| Asp | Gly | Arg | Ser | Leu 405 | Pro | Lys | Gly | Ile | Met 410 | Val | Leu | Leu | Ser | Ile 415 | Tyr |
| Gly | Leu | His | His | Asn | Pro | Lys | Val | Trp 425 | Pro | Asn | Pro | Glu | Val 430 | Phe | Asp |

Pro Phe Arg Phe Ala Pro Gly Ser Ala Gln His Ser His Ala Phe Leu Pro Phe Ser Gly Gly Ser Arg Asn Cys Ile Gly Lys Gln Phe Ala Met Asn Glu Leu Lys Val Ala Thr Ala Leu Thr Leu Leu Arg Phe Glu Leu Leu Pro Asp Pro Thr Arg Ile Pro Ile Pro Ile Ala Arg Leu Val Leu Lys Ser Lys Asn Gly Ile His Leu Arg Leu Arg Leu Pro Asn Pro Cys Glu Asp Lys Asp Gln Leu <210> 204 <211> 509 <212> PRT <213> Rattus norvegicus <400> 204 Met Ser Val Ser Ala Leu Ser Ser Thr Arg Phe Thr Gly Ser Ile Ser Gly Phe Leu Gln Val Ala Ser Val Leu Gly Leu Leu Leu Leu Val Lys Ala Val Gln Phe Tyr Leu Gln Arg Gln Trp Leu Leu Lys Ala Phe Gln Gln Phe Pro Ser Pro Pro Phe His Trp Phe Phe Gly His Lys Gln Phe Gln Gly Asp Lys Glu Leu Gln Gln Ile Met Thr Cys Val Glu Asn Phe Pro Ser Ala Phe Pro Arg Trp Phe Trp Gly Ser Lys Ala Tyr Leu Ile Val Tyr Asp Pro Asp Tyr Met Lys Val Ile Leu Gly Arg Ser Asp

Pro Lys Ala Asn Gly Val Tyr Arg Leu Leu Ala Pro Trp Ile Gly Tyr

Gly Leu Leu Leu Asn Gly Gln Pro Trp Phe Gln His Arg Arg Met Leu Thr Pro Ala Phe His Tyr Asp Ile Leu Lys Pro Tyr Val Lys Asn Met Ala Asp Ser Ile Arg Leu Met Leu Asp Lys Trp Glu Gln Leu Ala Gly Gln Asp Ser Ser Ile Glu Ile Phe Gln His Ile Ser Leu Met Thr Leu Asp Thr Val Met Lys Cys Ala Phe Ser His Asn Gly Ser Val Gln Val Asp Gly Asn Tyr Lys Ser Tyr Ile Gln Ala Ile Gly Asn Leu Asn Asp Leu Phe His Ser Arg Val Arg Asn Ile Phe His Gln Asn Asp Thr Ile Tyr Asn Phe Ser Ser Asn Gly His Leu Phe Asn Arg Ala Cys Gln Leu Ala His Asp His Thr Asp Gly Val Ile Lys Leu Arg Lys Asp Gln Leu Gln Asn Ala Gly Glu Leu Glu Lys Val Lys Lys Lys Arg Arg Leu Asp Phe Leu Asp Ile Leu Leu Leu Ala Arg Met Glu Asn Gly Asp Ser Leu Ser Asp Lys Asp Leu Arg Ala Glu Val Asp Thr Phe Met Phe Glu Gly His Asp Thr Thr Ala Ser Gly Val Ser Trp Ile Phe Tyr Ala Leu Ala Thr His Pro Glu His Gln Gln Arg Cys Arg Glu Glu Val Gln Ser Val Leu Gly Asp Gly Ser Ser Ile Thr Trp Asp His Leu Asp Gln Ile Pro Tyr Thr Thr Met Cys Ile Lys Glu Ala Leu Arg Leu Tyr Pro Pro

Val Pro Gly Ile Val Arq Glu Leu Ser Thr Ser Val Thr Phe Pro Asp 385 390 395 Gly Arg Ser Leu Pro Lys Gly Ile Gln Val Thr Leu Ser Ile Tyr Gly 405 410 415 Leu His His Asn Pro Lys Val Trp Pro Asn Pro Glu Val Phe Asp Pro 420 425 Ser Arg Phe Ala Pro Asp Ser Pro Arg His Ser His Ser Phe Leu Pro 440 435 Phe Ser Gly Gly Ala Arg Asn Cys Ile Gly Lys Gln Phe Ala Met Ser 455 460 Glu Met Lys Val Ile Val Ala Leu Thr Leu Leu Arg Phe Glu Leu Leu 470 475 Pro Asp Pro Thr Lys Val Pro Ile Pro Leu Pro Arg Leu Val Leu Lys 485 490 Ser Lys Asn Gly Ile Tyr Leu Tyr Leu Lys Lys Leu His 500 505 <210> 205 <211> 509 <212> PRT <213> Mus musculus <400> 205 Met Ser Val Ser Ala Leu Ser Pro Thr Arg Phe Ala Asp Ser Leu Ser 10 Gly Phe Leu Gln Val Ala Ser Val Leu Gly Leu Leu Leu Leu Val 20 25 Lys Ala Val Gln Phe Tyr Leu His Arg Gln Trp Leu Leu Lys Ala Phe Gln Gln Phe Pro Ser Pro Pro Phe His Trp Phe Phe Gly His Glu Lys 50 55 60 Phe Lys Gly Asp Gln Glu Leu Gln Glu Ile Val Ser Cys Ile Glu Asn 75

Phe Pro Ser Ala Phe Pro Arg Trp Phe Trp Gly Ser Lys Ala Tyr Leu

| Thr | Val | Tyr | Asp 100 | Pro | Asp | Tyr | Met | Lys 105 | Val | Ile | Leu | Gly | Arg 110 | Ser | Asp |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Pro | Lys | Ala 115 | Asn | Gly | Ala | Tyr | Arg 120 | Leu | Leu | Ala | Pro | Trp 125 | Ile | Gly | Tyr |
| Gly | Leu 130 | Leu | Leu | Leu | Asn | Gly 135 | Gln | Pro | Trp | Phe | Gln 140 | His | Arg | Ąrg | Met |
| Leu 145 | Thr | Pro | Ala | Phe | His 150 | Tyr | Asp | Ile | Leu | Lys 155 | Pro | Tyr | Val | Lys | Asn 160 |
| Met | Ala | Asp | Ser | Ile 165 | Arg | Leu | Met | Leu | Asp 170 | Lys | Trp | Glu | Arg | Leu 175 | Ala |
| Asp | Gln | Asp | Ser 180 | Ser | Ile | Glu | Ile | Phe 185 | Gln | His | Ile | Ser | Leu 190 | Met | Thr |
| Leu | Asp | Thr 195 | Val | Met | Lys | Cys | Ala 200 | Phe | Ser | His | Lys | Gly 205 | Ser | Val | Gln |
| Val | Asp 210 | Gly | Asn | Tyr | Arg | Thr 215 | Tyr | Leu | Gln | Ala | Ile 220 | Gly | Asp | Leu | Asn |
| Asn 225 | Leu | Phe | His | Ser | Arg 230 | Val | Arg | Asn | Ile | Phe 235 | His | Gln | Asn | Asp | Thr 240 |
| Ile | Tyr | Lys | Leu | Ser 245 | Ser | Asn | Gly | Arg | Leu 250 | Ala | Lys | Gln | Ala | Cys 255 | Gln |
| Leu | Ala | His | Asp 260 | His | Thr | Asp | Gly | Val 265 | Ile | Lys | Leu | Arg | Lys 270 | Asp | Gln |
| Leu | Gln | Asp 275 | Glu | Gly | Glu | Leu | Glu 280 | Lys | Ile | Lys | Lys | Lys 285 | Arg | Arg | Leu |
| Asp | Phe 290 | Leu | Asp | Ile | Leu | Leu 295 | Phe | Ala | Arg | Met | Glu 300 | Asn | Gly | Asp | Ser |
| Met 305 | Ser | Asp | Lys | Asp | Leu 310 | Arg | Ala | Glu | Val | Asp 315 | Thr | Phe | Met | Phe | Glu 320 |
| Gly | His | Asp | Thr | Thr 325 | Ala | Ser | Gly | Val | Ser 330 | Trp | Ile | Phe | Tyr | Ala 335 | Leu |

Ala Thr His Pro Asp His Gln Gln Arg Cys Arg Glu Glu Val Gln Ser

| 340 | 345 | 350 |
|-----|-----|-----|
| | | |

| Leu | Leu | Gly | Asp | Gly | Ser | Ser | Ile | Thr | Trp | Asp | His | Leu | Asp | GIn | Ile |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | 355 | | | | | 360 | | | | | 365 | | | |

Pro Tyr Thr Thr Met Cys Ile Lys Glu Ala Leu Arg Leu Tyr Pro Pro 370 380

Val Pro Gly Ile Val Arg Glu Leu Ser Thr Ser Val Thr Phe Pro Asp 385 395 400

Gly Arg Ser Leu Pro Lys Gly Val Gln Val Thr Leu Ser Ile Tyr Gly 405 410 415

Leu His His Asn Pro Lys Val Trp Pro Asn Pro Glu Val Phe Asp Pro 420 425 430

Ser Arg Phe Ala Pro Asp Ser Pro Arg His Ser His Ser Phe Leu Pro 435 440 445

Phe Ser Gly Gly Ala Arg Asn Cys Ile Gly Lys Gln Phe Ala Met Ser 450 455 460

Glu Leu Lys Val Ile Val Ala Leu Thr Leu Leu Arg Phe Glu Leu Leu 465 470 475 480

Pro Asp Pro Thr Arg Val Pro Met Pro Leu Ala Arg Leu Val Leu Lys 485 490 495

Ser Lys Asn Gly Ile Tyr Leu His Leu Lys Lys Leu His
500 505

<210> 206

<211> 509

<212> PRT

<213> Mus musculus

<400> 206

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Gly Phe Leu Gln Val Ala Ser Val Leu Gly Leu Leu Leu Leu Val 20 . 25 30

Lys Ala Val Gln Phe Tyr Leu His Arg Gln Trp Leu Leu Lys Ala Phe 35 40 45

| Gln | Gln 50 | Phe | Pro | Ser | Pro | Pro 55 | Phe | His | Trp | Phe | Phe 60 | Gly | His | Glu | Gln |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Phe 65 | Lys | Gly | Asp | His | Glu 70 | Leu | Gln | Glu | Ile | Val 75 | Ser | Cys | Ile | Glu | Asn 80 |
| Phe | Pro | Ser | Ala | Phe 85 | Pro | Arg | Trp | Phe | Trp 90 | Gly | Ser | Lys | Ala | Tyr 95 | Leu |
| Thr | Val | Tyr | Asp 100 | Pro | Asp | Tyr | Met | Lys 105 | Val | Ile | Leu | Gly | Arg 110 | Ser | Asp |
| Pro | Lys | Ala 115 | Asn | Gly | Ala | Tyr | Arg 120 | Leu | Leu | Ala | Pro | Trp 125 | Ile | Gly | Tyr |
| Gly | Leu 130 | Leu | Leu | Leu | Asn | Gly 135 | Gln | Pro | Trp | Phe | Gln 140 | His | Arg | Arg | Met |
| Leu 145 | Thr | Pro | Ala | Phe | His 150 | Tyr | Asp | Ile | Leu | Lys 155 | Pro | Tyr | Val | Lys | Asn 160 |
| Met | Ala | Asp | Ser | Ile 165 | Arg | Leu | Met | Leu | Asp 170 | Lys | Trp | Glu | Arg | Leu 175 | Ala |
| Asp | Gln | Asp | Ser 180 | Ser | Ile | Glu | Ile | Phe 185 | Gln | His | Ile | Ser | Leu 190 | Met | Thr |
| Leu | Asp | Thr 195 | Val | Met | Lys | Cys | Ala 200 | Phe | Ser | His | Lys | Gly 205 | Ser | Val | Gln |
| Val | Asp 210 | Gly | Asn | Tyr | Arg | Thr 215 | Tyr | Leu | Gln | Ala | Ile 220 | Gly | Asp | Leu | Asn |
| Asn 225 | Leu | Phe | His | Ser | Arg 230 | Val | Arg | Asn | Ile | Phe 235 | His | Gln | Asn | Asp | Thr 240 |
| Ile | Tyr | Lys | Leu | Ser 245 | Ser | Asn | Gly | Arg | Leu 250 | Ala | Lys | Gln | Ala | Cys 255 | Gln |
| Leu | Ala | His | Asp 260 | His | Thr | Asp | Gly | Val 265 | Ile | Lys | Leu | Arg | Lys 270 | Asp | Gln |
| Leu | Gln | Asp 275 | Glu | Gly | Glu | Leu | Glu 280 | Lys | Ile | Lys | Lys | Lys 285 | Arg | Arg | Leu |
| Asp | Phe 290 | Leu | Asp | Ile | Leu | Leu 295 | Phe | Ala | Arg | Met | Glu 300 | Asn | Gly | Asp | Ser |

Met Ser Asp Lys Asp Leu Arg Ala Glu Val Asp Thr Phe Met Phe Glu

315

320

Gly His Asp Thr Thr Ala Ser Gly Val Ser Trp Ile Phe Tyr Ala Leu 325 330 335

310

Ala Thr His Pro Asp His Gln Gln Arg Cys Arg Glu Glu Val Gln Ser 340 345 350

Leu Leu Gly Asp Gly Ser Ser Ile Thr Trp Asp His Leu Asp Gln Ile 355 360 365

Pro Tyr Thr Thr Met Cys Ile Lys Glu Ala Leu Arg Leu Tyr Pro Pro 370 380

Val Pro Gly Ile Val Arg Glu Leu Ser Thr Ser Val Thr Phe Pro Asp 385 390 395 400

Gly Arg Ser Leu Pro Lys Gly Val Gln Val Thr Leu Ser Ile Tyr Gly
405 410 415

Leu His His Asn Pro Lys Val Trp Pro Asn Pro Glu Val Phe Asp Pro 420 425 430

Ser Arg Phe Ala Pro Asp Ser Pro Arg His Ser His Ser Phe Leu Pro 435 440 445

Phe Ser Gly Gly Ala Arg Asn Cys Ile Gly Lys Gln Phe Ala Met Ser 450 455 460

Glu Leu Lys Val Ile Val Ala Leu Thr Leu Leu Arg Phe Glu Leu Leu 465 470 475 480

Pro Asp Pro Thr Arg Val Pro Met Pro Leu Ala Arg Leu Val Leu Lys
485 490 495

Ser Lys Asn Gly Ile Tyr Leu His Leu Lys Lys Leu His 500 505

<210> 207

305

<211> 519

<212> PRT

<213> Homo sapiens

<400> 207

Met Ser Val Ser Val Leu Ser Pro Ser Arg Leu Leu Gly Asp Val Ser

1 5 10 15

| Gly | Ile | Leu | Gln 20 | Ala | Ala | Ser | Leu | Leu 25 | Ile | Leu | Leu | Leu | Leu 30 | Leu | Ile |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Lys | Ala | Val 35 | Gln | Leu | Tyr | Leu | His 40 | Arg | Gln | Trp | Leu | Leu 45 | Lys | Ala | Leu |
| Gln | Gln 50 | Phe | Pro | Cys | Pro | Pro 55 | Ser | His | Trp | Leu | Phe 60 | Gly | His | Ile | Gln |
| Glu 65 | Leu | Gln | Gln | Asp | Gln 70 | Glu | Leu | Gln | Arg | Ile 75 | Gln | Lys | Trp | Val | Glu 80 |
| Thr | Phe | Pro | Ser | Ala 85 | Cys | Pro | His | Trp | Leu 90 | Trp | Gly | Gly | Lys | Val 95 | Arg |
| Val | Gln | Leu | Tyr 100 | Asp | Pro | Asp | Tyr | Met 105 | Lys | Val | Ile | Leu | Gly 110 | Arg | Ser |
| Asp | Pro | Lys 115 | Ser | His | Gly | Ser | Tyr 120 | Arg | Phe | Leu | Ala | Pro 125 | Trp | Ile | Gly |
| Tyr | Gly 130 | Leu | Leu | Leu | Leu | Asn 135 | Gly | Gln | Thr | Trp | Phe 140 | Gln | His | Arg | Arg |
| Met 145 | Leu | Thr | Pro | Ala | Phe 150 | His | Tyr | Asp | Ile | Leu 155 | Lys | Pro | Tyr | Val | Gly 160 |
| Leu | Met | Ala | Asp | Ser 165 | Val | Arg | Val | Met | Leu 170 | Asp | Lys | Trp | Glu | Glu 175 | Leu |
| Leu | Gly | Gln | Asp 180 | Ser | Pro | Leu | Glu | Val 185 | Phe | Gln | His | Val | Ser 190 | Leu | Met |
| Thr | Leu | Asp 195 | Thr | Ile | Met | Lys | Cys 200 | Ala | Phe | Ser | His | Gln 205 | Gly | Ser | Ile |
| Gln | Val 210 | Asp | Arg | Asn | Ser | Gln 215 | Ser | Tyr | Ile | Gln | Ala 220 | Ile | Ser | Asp | Leu |
| Asn 225 | Asn | Leu | Val | Phe | Ser 230 | Arg | Val | Arg | Asn | Ala 235 | Phe | His | Gln | Asn | Asp 240 |
| Thr | Ile | Tyr | Ser | Leu 245 | Thr | Ser | Ala | Gly | Arg 250 | Trp | Thr | His | Arg | Ala 255 | Cys |
| Gln | Leu | Ala | His 260 | Gln | His | Thr | Asp | Gln 265 | Val | Ile | Gln | Leu | Arg 270 | Lys | Ala |

Gln Leu Gln Lys Glu Gly Glu Leu Glu Lys Ile Lys Arq Lys Arq His Leu Asp Phe Leu Asp Ile Leu Leu Leu Ala Lys Met Glu Asn Gly Ser Ile Leu Ser Asp Lys Asp Leu Arg Ala Glu Val Asp Thr Phe Met Phe Glu Gly His Asp Thr Thr Ala Ser Gly Ile Ser Trp Ile Leu Tyr Ala Leu Ala Thr His Pro Lys His Gln Glu Arg Cys Arg Glu Glu Ile His Ser Leu Leu Gly Asp Gly Ala Ser Ile Thr Trp Asn His Leu Asp Gln Met Pro Tyr Thr Met Cys Ile Lys Glu Ala Leu Arg Leu Tyr Pro Pro Val Pro Gly Ile Gly Arg Glu Leu Ser Thr Pro Val Thr Phe Pro Asp Gly Arg Ser Leu Pro Lys Gly Ile Met Val Leu Leu Ser Ile Tyr Gly Leu His His Asn Pro Lys Val Trp Pro Asn Pro Glu Val Phe Asp Pro Ser Arg Phe Ala Pro Gly Ser Ala Gln His Ser His Ala Phe Leu Pro Phe Ser Gly Gly Ser Arg Asn Cys Ile Gly Lys Gln Phe Ala Met Asn Glu Leu Lys Val Ala Thr Ala Leu Thr Leu Leu Arg Phe Glu Leu Leu Pro Asp Pro Thr Arg Ile Pro Ile Pro Ile Ala Arg Leu Val Leu Lys Ser Lys Asn Gly Ile His Leu Arg Leu Arg Arg Leu Pro Asn Pro

Cys Glu Asp Lys Asp Gln Leu

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<210> 208
<211> 434
<212> PRT
<213> Homo sapiens
<400> 208
Pro Ala Pro Pro Thr His Trp Phe Leu Gly His Lys Leu Met Glu Lys
Tyr Pro Cys Ala Val Pro Leu Trp Val Gly Pro Phe Thr Met Phe Phe
             20
         35
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Ser Val His Asp Pro Asp Tyr Ala Lys Ile Leu Leu Lys Arg Gln Gly

Lys Asn Gln Glu Gly Phe Leu Pro Phe Ile Ser Gln Gly Lys Gly Leu

Ala Ala Leu Asp Gly Pro Lys Trp Phe Gln His Arg Arg Leu Leu Thr

Pro Gly Phe His Phe Asn Ile Leu Lys Ala Tyr Ile Glu Val Met Ala

His Ser Val Lys Met Met Leu Asn Lys Trp Glu Glu His Ile Ala Gln

Asn Ser Arg Leu Glu Leu Phe Gln His Val Ser Leu Met Thr Leu Asp

Ser Ile Met Lys Cys Ala Phe Ser His Gln Gly Ser Ile Gln Leu Asp

Arg Ser Ser Tyr Leu Lys Ala Val Phe Asn Leu Ser Lys Ile Ser Asn

Gln Arg Met Asn Asn Phe Leu His His Asn Asp Leu Val Phe Lys Phe

Ser Ser Gln Gly Gln Ile Phe Ser Lys Phe Asn Gln Glu Leu His Gln

His Leu Glu Lys Val Ile Gln Asp Arg Lys Glu Ser Leu Lys Asp Lys

Leu Lys Gln Asp Thr Thr Gln Lys Arg Arg Trp Asp Phe Leu Asp Ile

Ser Ser Ala Ile Ser Trp Ile Leu Tyr Cys Leu Ala Lys Tyr Pro Glu 260 265 270

His Gln Gln Arg Cys Arg Asp Glu Ile Arg Glu Leu Leu Gly Asp Gly 275 280 285

Ser Ser Ile Thr Trp His Leu Ser Gln Met Pro Tyr Thr Thr Met Cys 290 295 300

Ile Lys Glu Cys Leu Arg Leu Tyr Ala Pro Val Val Asn Ile Ser Arg 305 310 315 320

Leu Leu Asp Lys Pro Ile Thr Phe Pro Asp Gly Arg Ser Leu Pro Ala 325 330 335

Gly Ile Thr Val Val Leu Ser Ile Trp Gly Leu His His Asn Pro Ala 340 345 350

Val Trp Lys Asn Val Gln Val Phe Asp Pro Leu Arg Phe Ser Gln Glu 355 360 365

Asn Ser Asp Gln Arg His Pro Tyr Ala Tyr Leu Pro Phe Ser Ala Gly 370 380

Ser Arg Asn Cys Ile Gly Gln Glu Phe Ala Met Ile Glu Leu Lys Val 385 390 395 400

Thr Ile Ala Leu Ile Leu Leu His Phe Arg Val Thr Pro Asp Pro Thr 405 410 415

Arg Pro Leu Thr Phe Pro Asn His Phe Ile Leu Lys Pro Lys Asn Gly 420 425 430

Met Tyr

<210> 209

<211> 440

<212> PRT

| <213 | 3> A1 | tifi | icial | Sec | quenc | ce | | | | | | | | | |
|------------|---|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| | <220> <223> Description of Artificial Sequence: Cytochrome P450 domain sequence | | | | | | | | | | | | | | |
| <400 |)> 20 | 9 | | | | | | | | | | | | | |
| Pro 1 | Gly | Pro | Pro | Pro 5 | Leu | Pro | Leu | Ile | Gly 10 | Asn | Leu | Leu | Gln | Leu 15 | Gly |
| Arg | Gly | Pro | Ile 20 | His | Ser | Leu | Thr | Glu 25 | Leu | Arg | Lys | Lys | Tyr 30 | Gly | Pro |
| Val | Phe | Thr 35 | Leu | Tyr | Leu | Gly | Pro 40 | Arg | Pro | Val | Val | Val 45 | Val | Thr | Gly |
| Pro | Glu 50 | Ala | Val | Lys | Glu | Val 55 | Leu | Ile | Asp | Lys | Gly 60 | Glu | Glu | Phe | Ala |
| Gly 65 | Arg | Gly | Asp | Phe | Pro 70 | Val | Phe | Pro | Trp | Leu 75 | Gly | Tyr | Gly | Ile | Leu 80 |
| Phe | Ser | Asn | Gly | Pro 85 | Arg | Trp | Arg | Gln | Leu 90 | Arg | Arg | Leu | Leu | Thr 95 | Leu |
| Arg | Phe | Phe | Gly 100 | Met | Gly | Lys | Arg | Ser 105 | Lys | Leu | Glu | Glu | Arg 110 | Ile | Gln |
| Glu | Glu | Ala 115 | Arg | Asp | Leu | Val | Glu 120 | Arg | Leu | Arg | Lys | Glu 125 | Gln | Gly | Ser |
| Pro | Ile 130 | Asp | Ile | Thr | Glu | Leu 135 | Leu | Ala | Pro | Ala | Pro 140 | Leu | Asn | Val | Ile |
| Cys 145 | Ser | Leu | Leu | Phe | Gly 150 | Val | Arg | Phe | Asp | Tyr 155 | Glu | Asp | Pro | Glu | Phe 160 |
| Leu | Lys | Leu | Ile | Asp 165 | Lys | Leu | Asn | Glu | Leu 170 | Phe | Phe | Leu | Val | Ser 175 | Pro |
| Trp | Gly | Gln | Leu 180 | Leu | Asp | Phe | Phe | Arg 185 | Tyr | Leu | Pro | Gly | Ser 190 | His | Arg |
| Lys | Ala | Phe | Lys | Ala | Ala | Lys | Asp | Leu | Lys | Asp | Tyr | Leu | Asp | Lys | Leu |

Ile Glu Glu Arg Arg Glu Thr Leu Glu Pro Gly Asp Pro Arg Asp Phe

Leu Asp Ser Leu Leu Ile Glu Ala Lys Arg Glu Gly Gly Ser Glu Leu Thr Asp Glu Glu Leu Lys Ala Thr Val Leu Asp Leu Leu Phe Ala Gly Thr Asp Thr Thr Ser Ser Thr Leu Ser Trp Ala Leu Tyr Leu Leu Ala Lys His Pro Glu Val Gln Ala Lys Leu Arg Glu Glu Ile Asp Glu Val Ile Gly Arg Asp Arg Ser Pro Thr Tyr Asp Asp Arg Ala Asn Met Pro Tyr Leu Asp Ala Val Ile Lys Glu Thr Leu Arg Leu His Pro Val Val Pro Leu Leu Pro Arg Val Ala Thr Glu Asp Thr Glu Ile Asp Gly Tyr Leu Ile Pro Lys Gly Thr Leu Val Ile Val Asn Leu Tyr Ser Leu His Arg Asp Pro Lys Val Phe Pro Asn Pro Glu Glu Phe Asp Pro Glu Arg Phe Leu Asp Glu Asn Gly Lys Phe Lys Lys Ser Tyr Ala Phe Leu Pro Phe Gly Ala Gly Pro Arg Asn Cys Leu Gly Glu Arg Leu Ala Arg Met Glu Leu Phe Leu Phe Leu Ala Thr Leu Leu Gln Arg Phe Glu Leu Glu Leu Val Pro Pro Gly Asp Ile Pro Leu Thr Pro Lys Pro Leu Gly Leu Pro Ser Lys Pro Pro Leu Tyr

<210> 210

<211> 153

<212> PRT

<213> Mus musculus

<400> 210 Met Gly Ser Thr Met Glu Pro Pro Gly Gly Ala Tyr Leu His Leu Gly 10 Ala Val Thr Ser Pro Val Gly Thr Ala Arg Met Leu Gln Leu Ala Phe 20 25 Gly Cys Thr Thr Phe Ser Leu Val Ala His Arg Gly Gly Phe Gly Gly 35 40 Val Gln Gly Thr Phe Cys Met Ala Ala Trp Gly Phe Cys Phe Ala Phe 50 55 Ser Val Leu Val Val Ala Cys Glu Phe Thr Lys Leu His Ser Cys Leu 65 70 75 Arg Leu Ser Trp Gly Asn Phe Thr Ala Ala Phe Ala Met Leu Ala Thr 95 85 90 Leu Leu Cys Ala Thr Ala Ala Val Ile Tyr Pro Leu Tyr Phe Thr Arg 100 105 110 Leu Glu Cys Pro Pro Glu Pro Ala Gly Cys Met Val Ala Pro Cys Gln 115 120 Arg Pro Ala Pro Glu Ser Pro Trp Lys Asp Asp Asp Val Met Thr Ala 130 135 140 Met Glu Tyr Leu Ser Arg His Pro Thr 145 150

<210> 211 <211> 322

<212> PRT

<213> Homo sapiens

<400> 211

Met Pro Val Thr Val Thr Arg Thr Thr Ile Thr Thr Thr Thr Ser 1 5 10 15

Ser Ser Gly Leu Gly Ser Pro Met Ile Val Gly Ser Pro Arg Ala Leu 20 25 30

Thr Gln Pro Leu Gly Leu Leu Arg Leu Leu Gln Leu Val Ser Thr Cys
35 40 45

| Val | Ala 50 | Phe | Ser | Leu | Val | Ala 55 | Ser | Val | Gly | Ala | Trp 60 | Thr | Gly | Ser | Met |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly 65 | Asn | Trp | Ser | Met | Phe 70 | Thr | Trp | Cys | Phe | Cys 75 | Phe | Ser | Val | Thr | Leu 80 |
| Ile | Ile | Leu | Ile | Val 85 | Glu | Leu | Cys | Gly | Leu 90 | Gln | Ala | Arg | Phe | Pro 95 | Leu |
| Ser | Trp | Arg | Asn 100 | Phe | Pro | Ile | Thr | Phe 105 | Ala | Cys | Tyr | Ala | Ala 110 | Leu | Phe |
| Cys | Leu | Ser 115 | Ala | Ser | Ile | Ile | Tyr 120 | Pro | Thr | Thr | Tyr | Val 125 | Gln | Phe | Leu |
| Ser | His 130 | Gly | Arg | Ser | Arg | Asp 135 | His | Ala | Ile | Ala | Ala 140 | Thr | Phe | Phe | Ser |
| Cys 145 | Ile | Ala | Cys | Val | Ala 150 | Tyr | Ala | Thr | Glu | Val 155 | Ala | Trp | Thr | Arg | Ala 160 |
| Arg | Pro | Gly | Glu | Ile 165 | Thr | Gly | Tyr | Met | Ala 170 | Thr | Val | Pro | Gly | Leu 175 | Leu |
| Lys | Val | Leu | Glu 180 | Thr | Phe | Val | Ala | Cys 185 | Ile | Ile | Phe | Ala | Phe 190 | Ile | Ser |
| Asp | Pro | Asn 195 | Leu | Tyr | Gln | His | Gln 200 | Pro | Ala | Leu | Glu | Trp 205 | Cys | Val | Ala |
| Val | Tyr 210 | Ala | Ile | Cys | Phe | Ile 215 | Leu | Ala | Ala | Ile | Ala 220 | Ile | Leu | Leu | Asn |
| Leu 225 | Gly | Glu | Cys | Thr | Asn 230 | Val | Leu | Pro | Ile | Pro 235 | Phe | Pro | Ser | Phe | Leu 240 |
| Ser | Gly | Leu | Ala | Leu 245 | Leu | Ser | Val | Leu | Leu 250 | Tyr | Ala | Thr | Ala | Leu 255 | Val |
| Leu | Trp | Pro | Leu 260 | Tyr | Gln | Phe | Asp | Glu 265 | Lys | Tyr | Gly | Gly | Gln 270 | Pro | Arg |
| Arg | Ser | Arg 275 | Asp | Val | Ser | Cys | Ser 280 | Arg | Ser | His | Ala | Tyr 285 | Tyr | Val | Cys |
| Ala | Trp | Asp | Arg | Arg | Leu | Ala | Val | Ala | Ile | Leu | Thr | Ala | Ile | Asn | Leu |

Leu Ala Tyr Val Ala Asp Leu Val His Ser Ala His Leu Val Phe Val 305 310 315 320

<210> 212

Lys Val

<211> 296

<212> PRT

<213> Mus musculus

<400> 212

Met Pro Val Thr Val Thr Arg Thr Thr Ile Thr Thr Thr Thr Ser Ser 1 5 10 15

Ser Thr Thr Val Gly Ser Ala Arg Ala Leu Thr Gln Pro Leu Gly Leu 20 25 30

Leu Arg Leu Leu Gln Leu Ile Ser Thr Cys Val Ala Phe Ser Leu Val
35 40 45

Ala Ser Val Gly Ala Trp Thr Gly Pro Met Gly Asn Trp Ala Met Phe 50 55 60

Thr Trp Cys Phe Cys Phe Ala Val Thr Leu Ile Ile Leu Ile Val Glu 65 70 75 80

Leu Gly Gly Leu Gln Ala His Phe Pro Leu Ser Trp Arg Asn Phe Pro 85 90 95

Ile Thr Phe Ala Cys Tyr Ala Ala Leu Phe Cys Leu Ser Ser Ile 100 105 110

Ile Tyr Pro Thr Tyr Val Gln Phe Leu Ala His Gly Arg Thr Arg
115 120 125

Asp His Ala Ile Ala Ala Thr Thr Phe Ser Cys Val Ala Cys Leu Ala 130 135 140

Tyr Ala Thr Glu Val Ala Trp Thr Arg Ala Arg Pro Gly Glu Ile Thr 145 150 155 160

Gly Tyr Met Ala Thr Val Pro Gly Leu Leu Lys Val Phe Glu Thr Phe 165 170 175

Val Ala Cys Ile Ile Phe Ala Phe Ile Gly Glu Pro Leu Leu Tyr Asn 180 185 190

Gln Lys Pro Ala Leu Glu Trp Cys Val Ala Val Tyr Ala Ile Cys Phe 195 200 Ile Leu Ala Gly Val Thr Ile Leu Leu Asn Leu Gly Asp Cys Thr Asn 210 215 220 Val Leu Pro Ile Pro Phe Pro Thr Phe Leu Ser Gly Leu Ala Tyr Ser 225 230 235 Leu Phe Ser Phe Thr Pro Leu Pro Ser Ser Gly Pro Ser Thr Asn 245 250 Leu Ile Arg Asp Ile Arg Ala Asn Pro Ala Val Gln Trp Ile Gln Ala 260 265 270 Ala Leu Val Val Leu Val Ile Tyr Asn Pro Thr Arg Cys Val Ser Gly 275 280 Thr Asp Asp Trp Arg Cys Pro Ser 290 295 <210> 213 <211> 245 <212> PRT <213> Homo sapiens <400> 213 Met Thr Leu Val Ile Leu Leu Val Glu Leu Gly Gly Ser Gln Ala Arg 1 5 10 Phe Pro Leu Phe Trp Arg Asn Phe Pro Ile Thr Phe Ala Cys Tyr Ala 20 25 Ala Leu Leu Cys Leu Ser Ala Ser Ile Ile Tyr Pro Thr Tyr Leu 35 40 Gln Phe Leu Ser His Gly Arg Ser Arg Asp His Ala Ile Ala Ala Ile Val Phe Ser Gly Ile Ala Cys Val Ala Tyr Ala Thr Glu Val Thr Trp 65 70 75 Thr Arg Ala Arg Pro Gly Glu Ile Thr Asp Tyr Met Ala Ser Glu Leu

Gly Leu Leu Lys Val Leu Glu Thr Phe Val Ala Cys Leu Ile Phe Val

| 100 | 105 | 110 |
|-----|-----|-----|
| | | |

Phe Ile Asn Ser Pro Tyr Val Tyr His Asn Arg Pro Ala Leu Glu Trp 115 120 125

Cys Val Ala Val Tyr Ala Leu Cys Phe Val Leu Ala Ala Leu Thr Val 130 135 140

Leu Leu Ser Leu Gly His Cys Thr Asn Met Leu Pro Ile Arg Phe Pro 145 150 155 160

Ser Phe Leu Leu Gly Leu Ala Leu Leu Ser Val Leu Leu Tyr Ala Thr 165 170 175

Ala Leu Val Leu Trp Pro Leu Tyr Gln Phe Asn Glu Lys Tyr Gly Val 180 185 190

Gln Pro Trp Gln Thr Arg Asp Val Ser Cys Ser Asp Arg Asn Pro Tyr 195 200 205

Leu Val Cys Ile Trp Asp Arg Arg Leu Ala Val Thr Asn Leu Thr Ala 210 215 220

Val Asn Leu Leu Ala Tyr Val Gly Asp Leu Val Tyr Ser Ala His Leu 225 230 235 240

Val Phe Val Lys Val 245

<210> 214

<211> 331

<212> PRT

<213> Homo sapiens

<400> 214

Met Ala Arg Gln Arg Glu Glu Lys Arg Arg Thr Glu Gln Gly Phe Gly
1 5 10 15

Leu Lys Cys Ser Arg Leu Ile Ile Leu Pro Asn Ile Arg Ile Ile Tyr
20 25 30

Lys Phe Arg Ile Tyr Thr Cys Thr Leu Ser Glu Asn Thr Glu Asn Leu 35 40 45

Ala Leu Cys Ser Ser Asn Asn Gln Thr Lys Leu Asn Gln Thr Met Gln 50 55 60

and the second

| Met 65 | Leu | Lys | Pro | Asp | Leu 70 | Phe | Ser | Val | Ser | Ser 75 | Ser | Ala | Arg | Thr | Ala 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Met | Pro | Val | Thr 85 | Val | Thr | His | Pro | Thr 90 | Val | Thr | Thr | Thr | Met 95 | Arg |
| Ser | Pro | Thr | Val 100 | Val | Gly | Ser | Ser | Arg 105 | Ala | Leu | Ile | Gln | Pro 110 | Leu | Gly |
| Leu | Leu | Arg 115 | Leu | Leu | Gln | Leu | Val 120 | Ser | Thr | Cys | Val | Ala 125 | Leu | Ser | Leu |
| Val | Ala 130 | Ser | Cys | Phe | Cys | Phe 135 | Ala | Met | Thr | Leu | Val 140 | Ile | Leu | Leu | Val |
| Glu 145 | Leu | Gly | Gly | Ser | Gln 150 | Ala | Arg | Phe | Pro | Leu 155 | Phe | Trp | Arg | Asn | Phe 160 |
| Pro | Ile | Thr | Phe | Ala 165 | Cys | Tyr | Ala | Ala | Leu 170 | Leu | Cys | Leu | Ser | Ala 175 | Ser |
| Ile | Ile | Tyr | Pro 180 | Thr | Thr | Tyr | Leu | Gln 185 | Phe | Leu | Ser | His | Gly 190 | Arg | Ser |
| Arg | Asp | His 195 | Ala | Ile | Ala | Ala | Ile 200 | Val | Phe | Ser | Gly | Ile 205 | Ala | Cys | Val |
| Ala | Tyr 210 | Ala | Thr | Glu | Val | Thr 215 | Trp | Thr | Arg | Ala | Arg 220 | Pro | Gly | Glu | Ile |
| Thr 225 | Asp | Tyr | Met | Ala | Ser 230 | Glu | Leu | Gly | Leu | Leu 235 | Lys | Val | Leu | Glu | Thr 240 |
| Phe | Val | Ala | Cys | Leu 245 | Ile | Phe | Val | Phe | Ile 250 | Asn | Ser | Pro | Tyr | Val 255 | Tyr |
| His | Asn | Arg | Pro 260 | Ala | Leu | Glu | Trp | Trp 265 | Val | Ala | Val | Tyr | Ala 270 | Leu | Cys |
| Phe | Val | Leu 275 | Ala | Ala | Leu | Thr | Ile 280 | Leu | Leu | Ser | Leu | Gly 285 | His | Cys | Thr |
| Asn | Met 290 | Leu | Pro | Ile | Arg | Phe 295 | Pro | Ser | Phe | Leu | Leu 300 | Gly | Leu | Ala | Leu |
| Leu 305 | Ser | Val | Leu | Leu | Tyr 310 | Ala | Thr | Ala | Leu | Val 315 | Leu | Trp | Pro | Leu | Tyr 320 |

Gln Phe Asn Glu Asn Pro Gly Arg Arg Glu Met 325 330

<210> 215

<211> 365

<212> PRT

<213> Homo sapiens

<400> 215

Met Gly Tyr Cys Gln Gly Val Ser Gln Val Ala Val Val Leu Leu Met

1 5 10 15

Phe Pro Lys Glu Lys Glu Ala Phe Leu Ala Leu Ala Gln Leu Leu Thr 20 25 30

Ser Lys Asn Leu Pro Asp Thr Val Asp Gly Gln Leu Pro Met Gly Pro 35 40 45

His Ser Arg Ala Ser Gln Val Ala Pro Glu Thr Thr Ser Ser Lys Val 50 55 60

Asp Arg Gly Val Ser Thr Val Cys Gly Lys Pro Lys Val Val Gly Lys 65 70 75 80

Ile Tyr Gly Gly Arg Asp Ala Ala Gly Gln Trp Pro Trp Gln Ala 85 90 95

Ser Leu Leu Tyr Trp Gly Ser His Leu Cys Gly Ala Val Leu Ile Asp 100 105 110

Ser Cys Trp Leu Val Ser Thr Thr His Cys Phe Leu Lys Thr Ser Ser 115 120 125

Ser Phe Ile Leu Ser Ser Gly Arg Glu Phe Pro Gly Pro Cys Val Cys 130 135 140

Leu Leu Asn Pro Asp Met Arg Glu Ser Ile Gly Ser Val Cys Ala Gly
145 150 155 160

His Leu Gln Gly Phe Ser Ser Val Cys Thr Met Leu Leu Lys Ser Gln
165 170 175

Ala Pro Lys Asn Tyr Gln Val Leu Leu Gly Asn Ile Gln Leu Tyr His 180 185 190

Gln Thr Gln His Thr Gln Lys Met Ser Val His Arg Ile Ile Thr His 195 200 205

Pro Asp Phe Glu Lys Leu His Pro Phe Gly Ser Asp Ile Ala Met Leu 210 215 220 Gln Leu His Leu Pro Met Asn Phe Thr Ser Tyr Ile Val Pro Val Cys 230 235 225 Leu Pro Ser Arg Asp Met Gln Leu Pro Met Gln Leu Ser Pro Pro Phe 245 250 Tyr Leu Gln Glu Gly Lys Gly Asp Ser Gly Gly Pro Leu Val Cys Tyr 260 265 Leu Pro Ser Ala Trp Val Leu Val Gly Leu Ala Ser Trp Gly Leu Asp 280 285 Cys Arg His Pro Ala Tyr Pro Ser Ile Phe Thr Arg Val Thr Tyr Phe 295 Ile Asn Trp Ile Asp Lys Ile Met Arg Leu Thr Pro Leu Ser Asp Pro 310 315 Ala Leu Ala Pro His Thr Cys Ser Pro Pro Lys Pro Leu Arg Ala Ala 325 330 Gly Leu Pro Gly Pro Cys Ala Ala Leu Val Leu Pro Gln Thr Trp Leu 340 345 350 Leu Leu Pro Leu Thr Leu Arg Ala Pro Trp Gln Thr Leu 355 360 365 <210> 216 <211> 148 <212> PRT <213> Homo sapiens <400> 216 Cys Gly Lys Pro Lys Val Val Gly Lys Ile Tyr Gly Gly Arg Asp Ala Ala Ala Gly Gln Trp Pro Trp Gln Ala Ser Leu Leu Tyr Trp Gly Ser 20 25 His Leu Cys Gly Ala Val Leu Ile Asp Ser Cys Trp Leu Val Ser Thr 45

Thr His Cys Phe Leu Asn Lys Ser Gln Ala Pro Lys Asn Tyr Gln Val

50 55 60

Leu Leu Gly Asn Ile Gln Leu Tyr His Gln Thr Gln His Thr Gln Lys 65 70 75 80

Met Ser Val His Arg Ile Ile Thr His Pro Asp Phe Glu Lys Leu His 85 90 95

Pro Phe Gly Ser Asp Ile Ala Met Leu Gln Leu His Leu Pro Met Asn 100 105 110

Phe Thr Ser Tyr Ile Val Pro Val Cys Leu Pro Ser Arg Asp Met Gln 115 120 125

Leu Pro Ser Asn Val Ser Cys Trp Ile Thr Gly Trp Gly Met Ala Ile 130 135 140

Leu Gly Gly Leu 145

<210> 217

<211> 367

<212> PRT

<213> Mus musculus

<400> 217

Met Trp Gly Ser Arg Ala Gln Gln Ser Gly Pro Asp Arg Gly Gly Ala 1 5 10 15

Cys Leu Leu Ala Ala Phe Leu Leu Cys Phe Ser Leu Leu His Ala Gln
20 25 30

Asp Tyr Thr Pro Ser Gln Thr Pro Pro Pro Thr Ser Asn Thr Ser Leu
35 40 45

Lys Pro Arg Gly Arg Val Gln Lys Glu Leu Cys Gly Lys Thr Lys Phe 50 55 60

Gln Gly Lys Ile Tyr Gly Gly Gln Ile Ala Lys Ala Glu Arg Trp Pro 65 70 75 80

Trp Gln Ala Ser Leu Ile Phe Arg Gly Arg His Ile Cys Gly Ala Val $85 \hspace{1.5cm} 90 \hspace{1.5cm} 95$

Leu Ile Asp Lys Thr Trp Leu Leu Ser Ala Ala His Cys Phe Gln Arg
100 105 110

| Ser | Leu | Thr 115 | Pro | Ser | Asp | Tyr | Arg 120 | Ile | Leu | Leu | Gly | Tyr 125 | Asn | Gln | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser | Asn 130 | Pro | Ser | Asn | Tyr | Ser 135 | Arg | Gln | Met | Thr | Val 140 | Asn | Lys | Val | Ile |
| Leu 145 | His | Glu | Asp | Tyr | Ser 150 | Lys | Leu | Ser | Arg | Leu 155 | Glu | Lys | Asn | Ile | Val 160 |
| Leu | Ile | Gln | Leu | His 165 | His | Pro | Val | Ile | Tyr 170 | Ser | Thr | His | Ile | Phe 175 | Pro |
| Ala | Cys | Val | Pro 180 | Asp | Gly | Thr | Thr | Lys 185 | Val | Ser | Pro | Asn | Asn 190 | Leu | Cys |
| Trp | Ile | Ser 195 | Gly | Trp | Gly | Met | Leu 200 | Ser | Ala | Asp | Lys | Phe 205 | Leu | Gln | Ala |
| Pro | Phe 210 | Pro | Leu | Leu | Asp | Ala 215 | Glu | Val | Ser | Leu | Ile 220 | Asp | Glu | Glu | Glu |
| Cys 225 | Thr | Thr | Phe | Phe | Gln 230 | Thr | Pro | Glu | Val | Ser 235 | Ile | Thr | Glu | Tyr | Asp 240 |
| Val | Ile | Lys | Asp | Asp 245 | Val | Leu | Cys | Ala | Gly 250 | Asp | Leu | Thr | Asn | Gln 255 | Lys |
| Ser | Ser | Cys | Arg 260 | Gly | Asp | Ser | Gly | Gly 265 | Pro | Leu | Val | Cys | Phe 270 | Leu | Asn |
| Ser | Phe | Trp 275 | Tyr | Val | Val | Gly | Leu 280 | Ala | Asn | Trp | Asn | Gly 285 | Ala | Cys | Leu |
| Glu | Pro 290 | Ile | His | Ser | Pro | Asn 295 | Ile | Phe | Thr | Lys | Val 300 | Ser | Tyr | Phe | Ser |
| Asp 305 | Trp | Ile | Lys | Gln | Lys 310 | Lys | Ala | Asn | Thr | Pro 315 | Ala | Ala | Asp | Val | Ser 320 |
| Ser | Ala | Pro | Leu | Glu 325 | Glu | Met | Ala | Ser | Ser 330 | Leu | Arg | Gly | Trp | Gly 335 | Asn |
| Tyr | Ser | Ala | Gly 340 | Ile | Thr | Leu | Lys | Pro 345 | Arg | Ile | Ser | Thr | Thr 350 | Leu | Leu |
| Ser | Ser | Gln 355 | Ala | Leu | Leu | Leu | Gln 360 | Ser | Ile | Trp | Leu | Arg 365 | Ile | Leu | |

<210> 218 <211> 366 <212> PRT <213> Mus musculus <400> 218 Met Cys Gly Val Arg Ala Lys Lys Ser Gly Leu Ser Gly Tyr Gly Ala Gly Leu Leu Ala Ala Leu Leu Gly Val Ser Phe Leu Ser Gln His Ala Gln Thr Ala Glu Pro Thr Asn Val Thr Asn Ala Ala Asn Asn Thr Thr Ile Gln Ile Met Lys Ser Thr Leu Ser Leu Ser Glu Val Cys Gly Lys

Thr Lys Phe Gln Gly Lys Ile Tyr Gly Gln Ile Ala Gly Ala Glu Arg Trp Pro Trp Gln Ala Ser Leu Arg Leu Tyr Gly Arg His Ile Cys Gly Ala Val Leu Ile Asp Lys Asn Trp Val Leu Gly Ala Ala His Cys Phe Gln Arg Ser Gln Glu Pro Ser Asp Tyr His Val Met Leu Gly Tyr Thr Asp Leu Asn Ser Pro Thr Arg Tyr Ser Arg Thr Met Ser Val Gln Lys Val Ile Val His Lys Asp Tyr Asn Arg Phe His Thr Gln Gly Ser Asp Ile Val Leu Leu Gln Leu Arg Ser Ser Val Glu Tyr Ser Ser His Ile Leu Pro Ala Cys Val Pro Glu Glu Asn Ile Lys Ile Pro Lys Glu Lys Ala Cys Trp Ala Ser Gly Trp Gly Tyr Leu Arg Glu Asp Val Arg

Ile Pro Leu Pro Asn Glu Leu Tyr Glu Ala Glu Leu Ile Ile Met Ser

Asn Asp Gln Cys Lys Gly Phe Phe Pro Pro Pro Val Pro Gly Ser Ser 225 230 235 Arg Ser Tyr Tyr Ile Tyr Asp Asp Met Val Cys Ala Ala Asp Tyr Asp 245 250 Met Ser Lys Ser Ile Cys Ala Gly Asp Ser Gly Gly Pro Leu Val Cys 260 265 Leu Leu Glu Gly Ser Trp Tyr Val Val Gly Leu Thr Ser Trp Ser Ser 280 285 Thr Cys Glu Glu Pro Ile Val Ser Pro Ser Val Phe Ala Arg Val Ser 295 300 Tyr Phe Asp Lys Trp Ile Lys Asp Asn Lys Lys Ser Ser Ser Asn Ser 310 315 Lys Pro Gly Glu Ser Pro His His Pro Gly Ser Pro Glu Asn Glu Asn 325 330 Pro Glu Gly Asn Asn Lys Asn Gln Gly Thr Val Ile Lys Pro Val Cys 340 345 Thr Ala Leu Leu Ser Gln Thr Leu Leu Gln Gln Leu Ile 355 360 365 <210> 219 <211> 389 <212> PRT <213> Xenopus laevis <400> 219 Met Leu Gln Tyr Leu Ser Phe Val Leu Ile Phe Ile His His Gln Ala 10 Cys Gly Val Pro Val Ile Ser Asn Arg Ile Val Gly Gly Met Asp Ser 20 25 Lys Arg Gly Glu Trp Pro Trp Gln Ile Ser Leu Ser Tyr Lys Ser Asp Ser Ile Cys Gly Gly Ser Leu Leu Thr Asp Ser Trp Val Met Thr Ala Ala His Cys Ile Asp Ser Leu Asp Val Ser Tyr Tyr Thr Val Tyr Leu

- Gly Ala Tyr Gln Leu Ser Ala Pro Asp Asn Ser Thr Val Ser Arg Gly 85 90 95
- Val Lys Ser Ile Thr Lys His Pro Asp Phe Gln Tyr Glu Gly Ser Ser 100 105 110
- Gly Asp Ile Ala Leu Ile Glu Leu Glu Lys Pro Val Thr Phe Thr Pro 115 120 125
- Tyr Ile Leu Pro Ile Cys Leu Pro Ser Gln Asp Val Gln Phe Ala Ala 130 135 140
- Gly Thr Met Cys Trp Val Thr Gly Trp Gly Asn Ile Gln Glu Gly Thr 145 150 155 160
- Pro Leu Ile Ser Pro Lys Thr Ile Gln Lys Ala Glu Val Ala Ile Ile 165 170 175
- Asp Ser Ser Val Cys Gly Thr Met Tyr Glu Ser Ser Leu Gly Tyr Ile 180 185 190
- Pro Asp Phe Ser Phe Ile Gln Glu Asp Met Val Cys Ala Gly Tyr Lys
 195 200 205
- Glu Gly Arg Ile Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu Val 210 215 220
- Cys Asn Val Asn Asn Val Trp Leu Gln Leu Gly Ile Val Ser Trp Gly 225 230 235 240
- Tyr Gly Cys Ala Glu Pro Asn Arg Pro Gly Val Tyr Thr Lys Val Gln
 245 250 255
- Tyr Tyr Gln Asp Trp Leu Lys Thr Asn Val Pro Leu Ile Val Phe Ser 260 265 270
- Glu Glu Gly Pro Ser Val Ala Pro Ser Ile Gly Pro Ser Ile Ala Pro 275 280 285
- Ser Phe Gly Pro Ser Leu Gly Pro Arg Gly Val Ala Ser Thr Thr Ile 290 295 300
- Ser Gln Thr Glu Ala Gln Ser Val Asn Ser Ile Glu Ile Asp Lys Thr 305 310 315 320
- Asn Ser Thr Thr Ile Phe Glu Thr Glu Ala Met Ser Met Ser Asn Asn

325 330 335

Thr Thr Met Asn Glu Thr Phe Ser Leu Val Ser Ser Thr Ile Ser Thr 340 345 350

Ala Leu Arg Ile Asn Glu Thr Lys Thr Ile Asp Asn Glu Ala Gln Ile 355 360 365

His Ala Cys Ser Leu His Thr Ile Ala Leu Thr Leu Ile Tyr Leu Phe 370 380

Ile Arg Phe Phe Val 385

<210> 220

<211> 186

<212> PRT

<213> Homo sapiens

<400> 220

Lys Ile Tyr Gly Gly Arg Asp Ala Ala Ala Gly Gln Trp Pro Trp Gln
1 5 10 15

Ala Ser Leu Leu Tyr Trp Gly Ser His Leu Cys Gly Ala Val Leu Ile 20 25 30

Asp Ser Cys Trp Leu Val Ser Thr Thr His Cys Phe Lys Ser Gln Ala 35 40 45

Pro Lys Asn Tyr Gln Val Leu Leu Gly Asn Ile Gln Leu Tyr His Gln 50 55 60

Thr Gln His Thr Gln Lys Met Ser Val His Arg Ile Ile Thr His Pro 65 70 75 80

Asp Phe Glu Lys Leu His Pro Phe Gly Ser Asp Ile Ala Met Leu Gln 85 90 95

Leu His Leu Pro Met Asn Phe Thr Ser Tyr Ile Val Pro Val Cys Leu 100 105 110

Pro Ser Arg Asp Met Gln Leu Pro Ser Asn Val Ser Cys Trp Ile Thr
115 120 125

Gly Trp Gly Met Leu Thr Glu Asp Leu Cys Ser Gln Gly Asp Ser Gly 130 135 140

Ala Ser Trp Gly Leu Asp Cys Arg His Pro Ala Tyr Pro Ser Ile Phe 165 170 175

Thr Arg Val Thr Tyr Phe Ile Asn Trp Ile 180 185

<210> 221

<211> 230

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:Trypsin-like serine protease domain sequence

<400> 221

Arg Ile Val Gly Gly Ser Glu Ala Asn Ile Gly Ser Phe Pro Trp Gln
1 5 10 15

Val Ser Leu Gln Tyr Arg Gly Gly Arg His Phe Cys Gly Gly Ser Leu 20 25 30

Ile Ser Pro Arg Trp Val Leu Thr Ala Ala His Cys Val Tyr Gly Ser 35 40 45

Ala Pro Ser Ser Ile Arg Val Arg Leu Gly Ser His Asp Leu Ser Ser 50 55 60

Gly Glu Glu Thr Gln Thr Val Lys Val Ser Lys Val Ile Val His Pro 65 70 75 80

Asn Tyr Asn Pro Ser Thr Tyr Asp Asn Asp Ile Ala Leu Leu Lys Leu 85 90 95

Ser Glu Pro Val Thr Leu Ser Asp Thr Val Arg Pro Ile Cys Leu Pro 100 105 110

Ser Ser Gly Tyr Asn Val Pro Ala Gly Thr Thr Cys Thr Val Ser Gly 115 120 125

Trp Gly Arg Thr Ser Glu Ser Ser Gly Ser Leu Pro Asp Thr Leu Gln
130 135 140

Glu Val Asn Val Pro Ile Val Ser Asn Ala Thr Cys Arg Arg Ala Tyr

| | 145 | 150 | 155 | 160 |
|--|-----|-----|-----|-----|
|--|-----|-----|-----|-----|

Ser Gly Gly Pro Ala Ile Thr Asp Asn Met Leu Cys Ala Gly Gly Leu 165 170 175

Glu Gly Gly Lys Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu Val 180 185 190

Cys Asn Asp Pro Arg Trp Val Leu Val Gly Ile Val Ser Trp Gly Ser 195 200 205

Tyr Gly Cys Ala Arg Pro Asn Lys Pro Gly Val Tyr Thr Arg Val Ser 210 215 220

Ser Tyr Leu Asp Trp Ile 225 230

<210> 222

<211> 230

<212> PRT

<213> Homo sapiens

<400> 222

Arg Ile Val Gly Gly Ser Glu Ala Asn Ile Gly Ser Phe Pro Trp Gln
1 5 10 15

Val Ser Leu Gln Tyr Arg Gly Gly Arg His Phe Cys Gly Gly Ser Leu 20 25 30

Ile Ser Pro Arg Trp Val Leu Thr Ala Ala His Cys Val Tyr Gly Ser 35 40 45

Ala Pro Ser Ser Ile Arg Val Arg Leu Gly Ser His Asp Leu Ser Ser 50 55 60

Gly Glu Glu Thr Gln Thr Val Lys Val Ser Lys Val Ile Val His Pro 65 70 75 80

Asn Tyr Asn Pro Ser Thr Tyr Asp Asn Asp Ile Ala Leu Leu Lys Leu 85 90 95

Ser Glu Pro Val Thr Leu Ser Asp Thr Val Arg Pro Ile Cys Leu Pro 100 105 110

Ser Ser Gly Tyr Asn Val Pro Ala Gly Thr Thr Cys Thr Val Ser Gly
115 120 125

Trp Gly Arg Thr Ser Glu Ser Ser Gly Ser Leu Pro Asp Thr Leu Gln 135 140 130 Glu Val Asn Val Pro Ile Val Ser Asn Ala Thr Cys Arg Arg Ala Tyr 160 145 150 155 Ser Gly Gly Pro Ala Ile Thr Asp Asn Met Leu Cys Ala Gly Gly Leu 170 175 165 Glu Gly Gly Lys Asp Ala Cys Gln Gly Asp Ser Gly Gly Pro Leu Val 185 180 Cys Asn Asp Pro Arg Trp Val Leu Val Gly Ile Val Ser Trp Gly Ser 195 200 Tyr Gly Cys Ala Arg Pro Asn Lys Pro Gly Val Tyr Thr Arg Val Ser 220 210 215 Ser Tyr Leu Asp Trp Ile 225 230 <210> 223 <211> 217 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Trypsin-like serine protease domain sequence <400> 223 Ile Val Gly Gly Arg Glu Ala Gln Ala Gly Ser Phe Pro Trp Gln Val 10 Ser Leu Gln Val Ser Ser Gly His Phe Cys Gly Gly Ser Leu Ile Ser 20 25 Glu Asn Trp Val Leu Thr Ala Ala His Cys Val Ser Gly Ala Ser Ser 35 40 Val Arg Val Val Leu Gly Glu His Asn Leu Gly Thr Thr Glu Gly Thr 55 Glu Gln Lys Phe Asp Val Lys Lys Ile Ile Val His Pro Asn Tyr Asn 70 75

Pro Asp Thr Asn Asp Ile Ala Leu Leu Lys Leu Lys Ser Pro Val Thr

| 85 | 90 | 95 |
|----|----|----|
| | | |

Leu Gly Asp Thr Val Arg Pro Ile Cys Leu Pro Ser Ala Ser Ser Asp 100 105 110

Leu Pro Val Gly Thr Thr Cys Ser Val Ser Gly Trp Gly Arg Thr Lys
115 120 125

Asn Leu Gly Thr Ser Asp Thr Leu Gln Glu Val Val Val Pro Ile Val 130 135 140

Ser Arg Glu Thr Cys Arg Ser Ala Tyr Gly Gly Thr Val Thr Asp Thr 145 150 155 160

Met Ile Cys Ala Gly Ala Leu Gly Gly Lys Asp Ala Cys Gln Gly Asp 165 170 175

Ser Gly Gly Pro Leu Val Cys Ser Asp Gly Glu Leu Val Gly Ile Val 180 185 190

Ser Trp Gly Tyr Gly Cys Ala Val Gly Asn Tyr Pro Gly Val Tyr Thr 195 200 205

Arg Val Ser Arg Tyr Leu Asp Trp Ile 210 215

<210> 224

<211> 510

<212> PRT

<213> Homo sapiens

<400> 224

Met Asp Glu Lys Thr Lys Lys Ala Glu Glu Met Ala Leu Ser Leu Thr
1 5 10 15

Arg Ala Val Ala Gly Gly Asp Glu Gln Val Ala Met Lys Cys Ala Ile 20 25 30

Trp Leu Ala Glu Gln Arg Val Pro Leu Ser Val Gln Leu Lys Pro Glu 35 40 45

Val Ser Pro Thr Gln Asp Ile Arg Leu Trp Val Ser Val Glu Asp Ala 50 55 60

Gln Met His Thr Val Thr Ile Trp Leu Thr Val Arg Pro Asp Met Thr 65 70 75 80

| Val | Ala | Ser | Leu | Lys 85 | Asp | Met | Val | Phe | Leu 90 | Asp | Tyr | Gly | Phe | Pro 95 | Pro |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val | Leu | Gln | Gln 100 | Trp | Val | Ile | Gly | Gln 105 | Arg | Leu | Ala | Arg | Asp 110 | Gln | Glu |
| Thr | Leu | His 115 | Ser | His | Gly | Val | Arg 120 | Gln | Asn | Gly | Asp | Ser 125 | Ala | Tyr | Leu |
| Tyr | Leu 130 | Leu | Ser | Ala | Arg | Asn 135 | Thr | Ser | Leu | Asn | Pro 140 | Gln | Glu | Leu | Gln |
| Arg 145 | Glu | Arg | Gln | Leu | Arg 150 | Met | Leu | Glu | Asp | Leu 155 | Gly | Phe | Lys | Asp | Leu 160 |
| Thr | Leu | Gln | Pro | Arg 165 | Gly | Pro | Leu | Glu | Pro 170 | Gly | Pro | Pro | Lys | Pro 175 | Gly |
| Val | Pro | Gln | Glu 180 | Pro | Gly | Arg | Gly | Gln 185 | Pro | Asp | Ala | Val | Pro 190 | Glu | Pro |
| Pro | Pro | Val 195 | Gly | Trp | Gln | Cys | Pro 200 | Gly | Cys | Thr | Phe | Ile 205 | Asn | Lys | Pro |
| Thr | Arg 210 | Pro | Gly | Cys | Glu | Met 215 | Cys | Cys | Arg | Ala | Arg 220 | Pro | Glu | Ala | Tyr |
| Gln 225 | Val | Pro | Ala | Ser | Tyr 230 | Gln | Pro | Asp | Glu | Glu 235 | Glu | Arg | Ala | Arg | Leu 240 |
| Ala | Gly | Glu | Glu | Glu 245 | Ala | Leu | Arg | Gln | Tyr 250 | Gln | Gln | Arg | Lys | Gln 255 | Gln |
| Gln | Gln | Glu | Gly 260 | Asn | Tyr | Leu | Gln | His 265 | Val | Gln | Leu | Asp | Gln 270 | Arg | Ser |
| Leu | Val | Leu 275 | Asn | Thr | Glu | Pro | Ala 280 | Glu | Cys | Pro | Val | Cys 285 | Tyr | Ser | Val |
| Leu | Ala 290 | Pro | Gly | Glu | Ala | Val 295 | Val | Leu | Arg | Glu | Cys 300 | Leu | His | Thr | Phe |
| Cys 305 | Arg | Glu | Cys | Leu | Gln 310 | Gly | Thr | Ile | Arg | Asn 315 | Ser | Gln | Glu | Ala | Glu 320 |
| Val | Ser | Cys | Pro | Phe 325 | Ile | Asp | Asn | Thr | Tyr 330 | Ser | Cys | Ser | Gly | Lys 335 | Leu |

Phe Leu Asp Leu Gly Ile Ser Ile Ala Glu Asn Arg Ser Ala Phe Ser Tyr His Cys Lys Thr Pro Asp Cys Lys Gly Trp Cys Phe Phe Glu Asp Asp Val Asn Glu Phe Thr Cys Pro Val Cys Phe His Val Asn Cys Leu Leu Cys Lys Ala Ile His Glu Gln Met Asn Cys Lys Glu Tyr Gln Glu Asp Leu Ala Leu Arg Ala Gln Asn Asp Val Ala Ala Arg Gln Thr Thr Glu Met Leu Lys Val Met Leu Gln Gln Gly Glu Ala Met Arg Cys Pro Gln Cys Gln Ile Val Val Gln Lys Lys Asp Gly Cys Asp Trp Ile Arg Cys Thr Val Cys His Thr Glu Ile Cys Trp Val Thr Lys Gly Pro Arg Trp Gly Pro Gly Gly Pro Gly Asp Thr Ser Gly Gly Cys Arg Cys Arg Val Asn Gly Ile Pro Cys His Pro Ser Cys Gln Asn Cys His <210> 225 <211> 500 <212> PRT <213> Homo sapiens <400> 225 Met Ala Leu Ser Leu Thr Arg Ala Val Ala Gly Gly Asp Glu Gln Val Ala Met Lys Cys Ala Ile Trp Leu Ala Glu Gln Arg Val Pro Leu Ser Val Gln Leu Lys Pro Glu Val Ser Pro Thr Gln Asp Ile Arg Leu Trp

Leu Glu Arg Glu Ile Lys Ala Leu Leu Thr Pro Glu Asp Tyr Gln Arg

| Val | Ser 50 | Val | Glu | Asp | Ala | Gln 55 | Met | His | Thr | Val | Thr 60 | Ile | Trp | Leu | Thr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val 65 | Arg | Pro | Asp | Met | Thr 70 | Val | Ala | Ser | Leu | Lys 75 | Asp | Met | Val | Phe | Leu 80 |
| Asp | Tyr | Gly | Phe | Pro 85 | Pro | Val | Leu | Gln | Gln 90 | Trp | Val | Ile | Gly | Gln 95 | Arg |
| Leu | Ala | Arg | Asp 100 | Gln | Glu | Thr | Leu | His 105 | Ser | His | Gly | Val | Arg 110 | Gln | Asn |
| Gly | Asp | Ser 115 | Ala | Tyr | Leu | Tyr | Leu 120 | Leu | Ser | Ala | Arg | Asn 125 | Thr | Ser | Leu |
| Asn | Pro 130 | Gln | Glu | Leu | Gln | Arg 135 | Glu | Arg | Gln | Leu | Arg 140 | Met | Leu | Glu | Asp |
| Leu 145 | Gly | Phe | Lys | Asp | Leu 150 | Thr | Leu | Gln | Pro | Arg 155 | Gly | Pro | Leu | Glu | Pro 160 |
| Gly | Pro | Pro | Lys | Pro 165 | Gly | Val | Pro | Gln | Glu 170 | Pro | Gly | Arg | Gly | Gln 175 | Pro |
| Asp | Ala | Val | Pro 180 | Glu | Pro | Pro | Pro | Val 185 | Gly | Trp | Gln | Cys | Pro 190 | Gly | Cys |
| Thr | Phe | Ile 195 | Asn | Lys | Pro | Thr | Arg 200 | Pro | Gly | Cys | Glu | Met 205 | Суѕ | Cys | Arg |
| Ala | Arg 210 | Pro | Glu | Ala | Tyr | Gln 215 | Val | Pro | Ala | Ser | Tyr 220 | Gln | Pro | Asp | Glu |
| Glu 225 | Glu | Arg | Ala | Arg | Leu 230 | Ala | Gly | Glu | Glu | Glu 235 | Ala | Leu | Arg | Gln | Tyr 240 |
| Gln | Gln | Arg | Lys | Gln 245 | Gln | Gln | Gln | Glu | Gly 250 | Asn | Tyr | Leu | Gln | His 255 | Val |
| Gln | Leu | Asp | Gln 260 | Arg | Ser | Leu | Val | Leu 265 | Asn | Thr | Glu | Pro | Ala 270 | Glu | Cys |
| Pro | Val | Cys 275 | Tyr | Ser | Val | Leu | Ala 280 | Pro | Gly | Glu | Ala | Val 285 | Val | Leu | Arg |
| Glu | Cys 290 | Leu | His | Thr | Phe | Cys 295 | Arg | Glu | Cys | Leu | Gln 300 | Gly | Thr | Ile | Arg |

Asn Ser Gln Glu Ala Glu Val Ser Cys Pro Phe Ile Asp Asn Thr Tyr Ser Cys Ser Gly Lys Leu Leu Glu Arg Glu Ile Lys Ala Leu Leu Thr Pro Glu Asp Tyr Gln Arg Phe Leu Asp Leu Gly Ile Ser Ile Ala Glu Asn Arg Ser Ala Phe Ser Tyr His Cys Lys Thr Pro Asp Cys Lys Gly Trp Cys Phe Phe Glu Asp Asp Val Asn Glu Phe Thr Cys Pro Val Cys Phe His Val Asn Cys Leu Leu Cys Lys Ala Ile His Glu Gln Met Asn Cys Lys Glu Tyr Gln Glu Asp Leu Ala Leu Arg Ala Gln Asn Asp Val Ala Ala Arg Gln Thr Thr Glu Met Leu Lys Val Met Leu Gln Gln Gly Glu Ala Met Arg Cys Pro Gln Cys Gln Ile Val Val Gln Lys Lys Asp Gly Cys Asp Trp Ile Arg Cys Thr Val Cys His Thr Glu Ile Cys Trp Val Thr Lys Gly Pro Arg Trp Gly Pro Gly Gly Pro Gly Asp Thr Ser Gly Gly Cys Arg Cys Arg Val Asn Gly Ile Pro Cys His Pro Ser Cys Gln Asn Cys His

<210> 226

<211> 468

<212> PRT

<213> Homo sapiens

<400> 226

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| Val | Ser | Val | Glu 20 | Asp | Ala | Gln | Met | His 25 | Thr | Val | Thr | Ile | Trp | Leu | Thr |
| Val | Arg | Pro 35 | Asp | Met | Thr | Val | Ala 40 | Ser | Leu | Lys | Asp | Met 45 | Val | Phe | Leu · |
| Asp | Tyr 50 | Gly | Phe | Pro | Pro | Val 55 | Leu | Gln | Gln | Trp | Val 60 | Ile | Gly | Gln | Arg |
| Leu 65 | Ala | Arg | Asp | Gln | Glu 70 | Thr | Leu | His | Ser | His 75 | Gly | Val | Arg | Gln | Asn 80 |
| Gly | Asp | Ser | Ala | Tyr 85 | Leu | Tyr | Leu | Leu | Ser 90 | Ala | Arg | Asn | Thr | Ser 95 | Leu |
| Asn | Pro | Gln | Glu 100 | Leu | Gln | Arg | Glu | Arg 105 | Gln | Leu | Arg | Met | Leu 110 | Glu | Asp |
| Leu | Gly | Phe 115 | Lys | Asp | Leu | Thr | Leu 120 | Gln | Pro | Arg | Gly | Pro 125 | Leu | Glu | Pro |
| Gly | Pro 130 | Pro | Lys | Pro | Gly | Val 135 | Pro | Gln | Glu | Pro | Gly 140 | Arg | Gly | Gln | Pro |
| Asp 145 | Ala | Val | Pro | Glu | Pro 150 | Pro | Pro | Val | Gly | Trp 155 | Gln | Cys | Pro | Gly | Cys 160 |
| Thr | Phe | Ile | Asn | Lys 165 | Pro | Thr | Arg | Pro | Gly 170 | Cys | Glu | Met | Cys | Cys 175 | Arg |
| Ala | Arg | Pro | Glu 180 | Ala | Tyr | Gln | Val | Pro 185 | Ala | Ser | Tyr | Gln | Pro 190 | Asp | Glu |
| Glu | Glu | Arg 195 | Ala | Arg | Leu | Ala | Gly 200 | Glu | Glu | Glu | Ala | Leu 205 | Arg | Gln | Tyr |
| Gln | Gln 210 | Arg | Lys | Gln | Gln | Gln 215 | Gln | Glu | Gly | Asn | Tyr 220 | Leu | Gln | His | Val |
| Gln 225 | Leu | Asp | Gln | Arg | Ser 230 | Leu | Val | Leu | Asn | Thr 235 | Glu | Pro | Ala | Glu | Cys 240 |
| Pro | Val | Cys | Tyr | Ser 245 | Val | Leu | Ala | Pro | Gly 250 | Glu | Ala | Val | Val | Leu 255 | Arg |
| Glu | Cys | Leu | His | Thr | Phe | Cys | Arg | Glu | Cys | Leu | Gln | Gly | Thr | Ile | Arg |

Asn Ser Gln Glu Ala Glu Val Ser Cys Pro Phe Ile Asp Asn Thr Tyr 275 280 285

Ser Cys Ser Gly Lys Leu Leu Glu Arg Glu Ile Lys Ala Leu Leu Thr 290 295 300

Pro Glu Asp Tyr Gln Arg Phe Leu Asp Leu Gly Ile Ser Ile Ala Glu 305 310 315 320

Asn Arg Ser Ala Phe Ser Tyr His Cys Lys Thr Pro Asp Cys Lys Gly 325 330 335

Trp Cys Phe Phe Glu Asp Asp Val Asn Glu Phe Thr Cys Pro Val Cys 340 345 350

Phe His Val Asn Cys Leu Leu Cys Lys Ala Ile His Glu Gln Met Asn 355 360 365

Cys Lys Glu Tyr Gln Glu Asp Leu Ala Leu Arg Ala Gln Asn Asp Val 370 375 380

Ala Ala Arg Gln Thr Thr Glu Met Leu Lys Val Met Leu Gln Gln Gly 385 390 395 400

Glu Ala Met Arg Cys Pro Gln Cys Gln Ile Val Val Gln Lys Lys Asp 405 410 415

Gly Cys Asp Trp Ile Arg Cys Thr Val Cys His Thr Glu Ile Cys Trp
420 425 430

Val Thr Lys Gly Pro Arg Trp Gly Pro Gly Gly Pro Gly Asp Thr Ser 435 440 445

Gly Gly Cys Arg Cys Arg Val Asn Gly Ile Pro Cys His Pro Ser Cys 450 455 460

Gln Asn Cys His 465

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<212> PRT

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<400> 227

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| Ala | Ile | Lys | Tyr 20 | Ala | Thr | Trp | Leu | Ala 25 | Glu | Gln | Arg | Val | Pro 30 | Leu | Arg |
| Val | Gln | Val 35 | Lys | Pro | Glu | Val | Ser 40 | Pro | Thr | Gln | Asp | Ile 45 | Arg | Leu | Cys |
| Val | Ser 50 | Val | Glu | Asp | Ala | Tyr 55 | Met | His | Thr | Val | Thr 60 | Ile | Trp | Leu | Thr |
| Val 65 | Arg | Pro | Asp | Met | Thr 70 | Val | Ala | Ser | Leu | Lys 75 | Asp | Met | Val | Phe | Leu 80 |
| Asp | Tyr | Gly | Phe | Pro 85 | Pro | Ser | Leu | Gln | Gln 90 | Trp | Val | Val | Gly | Gln 95 | Arg |
| Leu | Ala | Arg | Asp 100 | Gln | Glu | Thr | Leu | His 105 | Ser | His | Gly | Ile | Arg 110 | Arg | Asn |
| Gly | Asp | Gly 115 | Ala | Tyr | Leu | Tyr | Leu 120 | Leu | Ser | Ala | Arg | Asn 125 | Thr | Ser | Leu |
| Asn | Pro 130 | Gln | Glu | Leu | Gln | Arg 135 | Gln | Arg | Gln | Leu | Arg 140 | Met | Leu | Glu | Asp |
| Leu 145 | Gly | Phe | Lys | Asp | Leu 150 | Thr | Leu | Gln | Ser | Arg 155 | Gly | Pro | Leu | Glu | Pro 160 |
| Val | Leu | Pro | Lys | Pro 165 | Arg | Thr | Asn | Gln | Glu 170 | Pro | Gly | Gln | Pro | Asp 175 | Ala |
| Ala | Pro | Glu | Ser 180 | Pro | Pro | Val | Gly | Trp 185 | Gln | Cys | Pro | Gly | Cys 190 | Thr | Phe |
| Ile | Asn | Lys 195 | Pro | Thr | Arg | Pro | Gly 200 | Cys | Glu | Met | Cys | Cys 205 | Arg | Ala | Arg |
| Pro | Glu 210 | Thr | Tyr | Gln | Ile | Pro 215 | Ala | Ser | Tyr | Gln | Pro 220 | Asp | Glu | Glu | Glu |
| Arg 225 | Ala | Arg | Leu | Ala | Gly 230 | Glu | Glu | Glu | Ala | Leu 235 | Arg | Gln | Tyr | Gln | Gln 240 |
| Arg | Lys | Gln | Gln | Gln 245 | Gln | Glu | Gly | Asn | Tyr 250 | | Gln | His | Val | Gln 255 | Leu |

| Glu | Gln | Arg | Ser 260 | Leu | Val | Leu | Asn | Thr 265 | Glu | Pro | Thr | Glu | Cys 270 | Pro | Val |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Cys | Tyr | Ser 275 | Val | Leu | Ala | Pro | Gly 280 | Glu | Ala | Val | Val | Leu 285 | Arg | Glu | Cys |
| Leu | His 290 | Thr | Phe | Cys | Arg | Glu 295 | Cys | Leu | Gln | Gly | Thr 300 | Ile | Arg | Asn | Ser |
| Gln 305 | Glu | Ala | Glu | Val | Ala 310 | Cys | Pro | Phe | Ile | Asp 315 | Ser | Thr | Tyr | Ser | Cys 320 |
| Pro | Gly | Lys | Leu | Leu 325 | Glu | Arg | Glu | Ile | Arg 330 | Ala | Leu | Leu | Ser | Pro 335 | Glu |
| Asp | Tyr | Gln | Arg 340 | Phe | Leu | Asp | Leu | Gly 345 | Val | Ser | Ile | Ala | Glu 350 | Asn | Arg |
| Ser | Thr | Leu 355 | Ser | Tyr | His | Cys | Lys 360 | Thr | Pro | Asp | Cys | Arg 365 | Gly | Trp | Cys |
| Phe | Phe 370 | Glu | Asp | Asp | Val | Asn 375 | Glu | Phe | Thr | Cys | Pro 380 | Val | Cys | Thr | Arg |
| Val 385 | Asn | Cys | Leu | Leu | Cys 390 | Lys | Ala | Ile | His | Glu 395 | His | Met | Asn | Cys | Arg 400 |
| Glu | Tyr | Gln | Asp | Asp 405 | Leu | Ala | Leu | Arg | Ala 410 | Gln | Asn | Asp | Val | Ala 415 | Ala |
| Arg | Gln | Thr | Thr 420 | Glu | Met | Leu | Lys | Val 425 | Met | Leu | Gln | Gln | Gly 430 | Glu | Ala |
| Met | His | Cys 435 | Pro | Gln | Cys | Arg | Ile 440 | Val | Val | Gln | Lys | Lys 445 | Asp | Gly | Cys |
| Asp | Trp 450 | Ile | Arg | Cys | Thr | Val 455 | Cys | His | Thr | Glu | Ile 460 | Cys | Trp | Val | Thr |
| Lys 465 | Gly | Pro | Arg | Trp | Gly 470 | Pro | Gly | Gly | Pro | Gly 475 | Asp | Thr | Ser | Gly | Gly 480 |
| Cys | Arg | Cys | Arg | Val 485 | Asn | Gly | Ile | Pro | Cys 490 | His | Pro | Ser | Cys | Gln 495 | Asn |

Cys His

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Pro Glu Ala Tyr Gln Ile Pro Ala Ser Tyr Gln Pro Asp Glu Glu Glu

| Arg 225 | Ala | Arg | Leu | Ala | Gly 230 | Glu | Glu | Glu | Ala | Leu 235 | Arg | Gln | Tyr | Glu | Gln 240 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Arg | Lys | Gln | Gln | Gln 245 | Gln | Glu | Gly | Asn | Tyr 250 | Leu | Gln | His | Val | Gln 255 | Leu |
| Glu | Gln | Arg | Ser 260 | Leu | Val | Leu | Asn | Thr 265 | Glu | Pro | Ala | Glu | Cys 270 | Pro | Val |
| Cys | Tyr | Ser 275 | Val | Leu | Ala | Pro | Gly 280 | Glu | Ala | Val | Val | Leu 285 | Arg | Glu | Cys |
| Leu | His 290 | Thr | Phe | Cys | Arg | Glu 295 | Cys | Leu | Gln | Gly | Thr 300 | Ile | Arg | Asn | Ser |
| Gln 305 | Glu | Ala | Glu | Val | Ser 310 | Cys | Pro | Phe | Ile | Asp 315 | Asn | Thr | Tyr | Ser | Cys 320 |
| Pro | Gly | Lys | Leu | Leu 325 | Glu | Arg | Glu | Ile | Arg 330 | Ala | Leu | Leu | Ser | Pro 335 | Glu |
| Asp | Tyr | Gln | Arg 340 | Phe | Leu | Asp | Leu | Gly 345 | Val | Ser | Ile | Ala | Glu 350 | Asn | Arg |
| Ser | Thr | Leu 355 | Ser | Tyr | His | Cys | Lys 360 | Thr | Pro | Asp | Cys | Arg 365 | Gly | Trp | Cys |
| Phe | Phe 370 | Glu | Asp | Asp | Val | Asn 375 | Glu | Phe | Thr | Cys | Pro 380 | Val | Cys | Thr | Arg |
| Val 385 | Asn | Cys | Leu | Leu | Cys 390 | Lys | Ala | Ile | His | Glu 395 | Arg | Met | Asn | Cys | Arg 400 |
| Glu | Tyr | Gln | Asp | Asp 405 | Leu | Ala | His | Arg | Ala 410 | Arg | Asn | Asp | Val | Ala 415 | Ala |
| Gln | Gln | Thr | Thr 420 | Glu | Met | Leu | Arg | Val 425 | Met | Leu | Gln | Gln | Gly 430 | Glu | Ala |
| Met | Tyr | Cys 435 | Pro | Gln | Cys | Arg | Ile 440 | Val | Val | Gln | Lys | Lys 445 | Asp | Gly | Cys |
| Asp | Trp 450 | Ile | Arg | Cys | Thr | Val 455 | Cys | His | Thr | Glu | Ile 460 | Cys | Trp | Val | Thr |
| Lys 465 | Gly | Pro | Arg | Trp | Gly 470 | Pro | Gly | Gly | Pro | Gly 475 | Asp | Thr | Ser | Gly | Gly 480 |

485 490 Cys His <210> 229 <211> 32 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: zf-RanBP domain sequence <400> 229 Arg Ala Gly Ser Asp Trp Asp Cys Ile Ser Ser Cys Leu Val Gln Asn 10 Phe Ala Thr Ser Thr Lys Cys Val Ala Cys Gln Ala Pro Lys Pro Ser 25 20 <210> 230 <211> 29 <212> PRT <213> Homo sapiens <400> 230 Pro Val Gly Trp Gln Cys Pro Gly Cys Thr Phe Ile Asn Lys Pro Thr 5 10 15 1 Arg Pro Gly Cys Glu Met Cys Cys Arg Ala Arg Pro Glu 25 20 <210> 231 <211> 53 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: zf-C3HC4

Cys Arg Cys Arg Val Asn Gly Ile Pro Cys His Pro Ser Cys Gln Asn

domain sequence

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<210> 234 <211> 8

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<223> Description of Artificial Sequence: IBR domain
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Asp Leu Lys Trp Cys Pro Gly Pro Asp Cys Ser Tyr Ala Val Arg Leu
             20
                                  25
Thr Glu Val Ser Ser Ser Thr Glu Leu Ala Glu Pro Pro Arg Val Glu
                              40
Cys Lys Lys Pro Ala Cys Gly Thr Ser Phe Cys Phe Lys Cys Gly Ala
                          55
Glu Trp His Ala Pro Val Ser Cys
 65
                     70
<210> 236
<211> 61
<212> PRT
<213> Homo sapiens
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Gln Arg Phe Leu Asp Leu Gly Ile Ser Ile Ala Glu Asn Arg Ser Ala
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                  5
                                      10
                                                           15
Phe Ser Tyr His Cys Lys Thr Pro Asp Cys Lys Gly Trp Cys Phe Phe
                                                       30
             20
                                  25
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35

Glu Asp Asp Val Asn Glu Phe Thr Cys Pro Val Cys Phe His Val Asn

Cys Leu Leu Cys Lys Ala Ile His Glu Gln Met Asn Cys 55 <210> 237 <211> 61 <212> PRT <213> Homo sapiens <400> 237 Thr Ile Trp Leu Thr Val Arg Pro Asp Met Thr Val Ala Ser Leu Lys 1 5 10 15 Asp Met Val Phe Leu Asp Tyr Gly Phe Pro Pro Val Leu Gln Gln Trp 20 25 Val Ile Gly Gln Arg Leu Ala Arg Asp Gln Glu Thr Leu His Ser His 35 40 Gly Val Arg Gln Asn Gly Asp Ser Ala Tyr Leu Tyr Leu 55 50 60 <210> 238 <211> 60 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Ubiquitin homologues domain sequence Thr Ile Thr Leu Glu Val Lys Pro Ser Asp Thr Val Ser Glu Leu Lys 1 5 10 15 Glu Lys Ile Ala Asp Leu Glu Gly Ile Pro Pro Glu Gln Gln Arg Leu 20 25 Ile Tyr Lys Gly Lys Val Leu Glu Asp Asp Arg Thr Leu Ala Glu Tyr 35 40 45

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Gly Ile Gln Asp Gly Ser Thr Ile His Leu Val Leu

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<213> Homo sapiens

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Met Asn Pro Glu Ser Ser Ile Phe Ile Glu Asp Tyr Leu Lys Tyr Phe 1 5 10 15

Gln Asp Gln Val Ser Arg Glu Asn Leu Leu Gln Leu Leu Thr Asp Asp 20 25 30

Glu Ala Trp Asn Gly Phe Val Ala Ala Ala Glu Leu Pro Arg Asp Glu 35 40 45

Ala Asp Glu Leu Arg Lys Ala Leu Asn Lys Leu Ala Ser His Met Val 50 55 60

Met Lys Asp Lys Asn Arg His Asp Lys Asp Gln Gln His Arg Gln Trp 65 70 75 80

Phe Leu Lys Glu Phe Pro Arg Leu Lys Arg Glu Leu Glu Asp His Ile 85 90 95

Arg Lys Leu Arg Ala Leu Ala Glu Glu Val Glu Gln Val His Arg Gly
100 105 110

Thr Thr Ile Ala Asn Val Val Ser Asn Ser Val Gly Thr Thr Ser Gly 115 120 125

Ile Leu Thr Leu Leu Gly Leu Gly Leu Ala Pro Phe Thr Glu Gly Ile 130 135 140

Ser Phe Val Leu Leu Asp Thr Gly Met Gly Leu Gly Ala Ala Ala 145 150 155 160

Val Ala Gly Ile Thr Cys Ser Val Val Glu Leu Val Asn Lys Leu Arg 165 170 175

Ala Arg Ala Gln Ala Arg Asn Leu Asp Gln Ser Gly Thr Asn Val Ala 180 185 190

Lys Val Met Lys Glu Phe Val Gly Gly Asn Thr Pro Asn Val Leu Thr 195 200 205

Leu Val Asp Asn Trp Tyr Gln Val Thr Gln Gly Ile Gly Arg Asn Ile 210 215 220

Arg Ala Ile Arg Arg Ala Arg Ala Asn Pro Gln Leu Gly Ala Tyr Ala

Pro Pro Pro His Val Ile Gly Arg Ile Ser Ala Glu Gly Glu Gln 245 250 255

Val Glu Arg Val Val Glu Gly Pro Ala Gln Ala Met Ser Arg Gly Thr 260 265 270

Met Ile Val Gly Ala Ala Thr Gly Gly Ile Leu Leu Leu Leu Asp Val 275 280 285

Val Ser Leu Ala Tyr Glu Ser Lys His Leu Leu Glu Gly Ala Lys Ser 290 295 300

Glu Ser Ala Glu Glu Leu Lys Lys Arg Ala Gln Glu Leu Glu Gly Lys 305 310 315 320

Leu Asn Phe Leu Thr Lys Ile His Glu Met Leu Gln Pro Gly Gln Asp 325 330 335

Gln

<210> 240

<211> 337

<212> PRT

<213> Homo sapiens

<400> 240

Met Asn Pro Glu Ser Ser Ile Phe Ile Glu Asp Tyr Leu Lys Tyr Phe 1 5 10 15

Gln Asp Gln Val Ser Arg Glu Asn Leu Leu Gln Leu Leu Thr Asp Asp 20 25 30

Glu Ala Trp Asn Gly Phe Val Ala Ala Ala Glu Leu Pro Arg Asp Glu
35 40 45

Ala Asp Glu Leu Arg Lys Ala Leu Asn Lys Leu Ala Ser His Met Val 50 55 60

Met Lys Asp Lys Asn Arg His Asp Lys Asp Gln Gln His Arg Gln Trp 65 70 75 80

Phe Leu Lys Glu Phe Pro Arg Leu Lys Arg Glu Leu Glu Asp His Ile 85 90 95

| Arg | Lys | Leu | Arg 100 | Ala | Leu | Ala | Glu | Glu 105 | Val | Glu | Gln | Val | His 110 | Arg | Gly |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Thr | Thr | Ile 115 | Ala | Asn | Val | Val | Ser 120 | Asn | Ser | Val | Gly | Thr 125 | Thr | Ser | Gly |
| Ile | | Thr | Leu | Leu | Gly | Leu 135 | Gly | Leu | Ala | Pro | Phe 140 | Thr | Glu | Gly | Ile |
| Ser 145 | Phe | Val | Leu | Leu | Asp 150 | Thr | Gly | Met | Gly | Leu 155 | Gly | Ala | Ala | Ala | Ala 160 |
| Val | Ala | Gly | Ile | Thr 165 | Cys | Ser | Val | Val | Glu 170 | Leu | Val | Asn | Lys | Leu 175 | Arg |
| Ala | Arg | Ala | Gln 180 | Ala | Arg | Asn | Leu | Asp 185 | Gln | Ser | Gly | Thr | Asn 190 | Val | Ala |
| Lys | Val | Met 195 | Lys | Glu | Phe | Val | Gly 200 | Gly | Asn | Thr | Pro | Asn 205 | Val | Leu | Thr |
| Leu | Val 210 | Asp | Asn | Trp | Tyr | Gln 215 | Val | Thr | Gln | Gly | Ile 220 | Gly | Arg | Asn | Ile |
| Arg 225 | Ala | Ile | Arg | Arg | Ala 230 | Arg | Ala | Asn | Pro | Gln 235 | Leu | Gly | Ala | Tyr | Ala 240 |
| Pro | Pro | Pro | His | Ile 245 | Ile | Gly | Arg | Ile | Ser 250 | Ala | Glu | Gly | Gly | Glu 255 | Gln |
| Val | Glu | Arg | Val 260 | Val | Glu | Gly | Pro | Ala 265 | Gln | Ala | Met | Ser | Arg 270 | Gly | Thr |
| Met | Ile | Val 275 | Gly | Ala | Ala | Thr | Gly 280 | Gly | Ile | Leu | Leu | Leu 285 | Leu | Asp | Val |
| Val | Ser 290 | Leu | Ala | Tyr | Glu | Ser 295 | Lys | His | Leu | Leu | Glu 300 | Gly | Ala | Lys | Ser |
| Glu 305 | Ser | Ala | Glu | Glu | Leu 310 | Lys | Lys | Arg | Ala | Gln 315 | Glu | Leu | Glu | Gly | Lys 320 |
| Leu | Asn | Phe | Leu | Thr 325 | Lys | Ile | His | Glu | Met 330 | Leu | Gln | Pro | Gly | Gln 335 | Asp |

Gln

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| Leu 1 | Ala | Ser | His | Met 5 | Val | Met | Lys | Asp | Lys 10 | Asn | Arg | His | Asp | Lys 15 | Asp |
| Gln | Gln | His | Arg 20 | Gln | Trp | Phe | Leu | Lys 25 | Glu | Phe | Pro | Arg | Leu 30 | Lys | Arg |
| Glu | Leu | Glu 35 | Asp | His | Ile | Arg | Lys 40 | Leu | Arg | Ala | Leu | Ala 45 | Glu | Glu | Val |
| Glu | Gln 50 | Val | His | Arg | Gly | Thr 55 | Thr | Ile | Ala | Asn | Val 60 | Val | Ser | Asn | Ser |
| Val 65 | Gly | Thr | Thr | Ser | Gly 70 | Ile | Leu | Thr | Leu | Leu 75 | Gly | Leu | Gly | Leu | Ala 80 |
| Pro | Phe | Thr | Glu | Gly 85 | Ile | Ser | Phe | Val | Leu 90 | Leu | Asp | Thr | Gly | Met 95 | Gly |
| Leu | Gly | Ala | Ala 100 | Ala | Ala | Val | Ala | Gly 105 | Ile | Thr | Cys | Ser | Val 110 | Val | Glu |
| Leu | Val | Asn 115 | Lys | Leu | Arg | Ala | Arg 120 | Ala | Gln | Ala | Arg | Asn 125 | Leu | Asp | Gln |
| Ser | Gly 130 | Thr | Asn | Val | Ala | Lys 135 | Val | Met | Lys | Glu | Phe 140 | Val | Gly | Gly | Asn |
| Thr 145 | Pro | Asn | Val | Leu | Thr 150 | Leu | Val | Asp | Asn | Trp 155 | Tyr | Gln | Val | Thr | Gln 160 |
| Gly | Ile | Gly | Arg | Asn 165 | Ile | Arg | Ala | Ile | Arg 170 | Arg | Ala | Arg | Ala | Asn 175 | Pro |
| Gln | Leu | Gly | Ala 180 | Tyr | Ala | Pro | Pro | Pro 185 | His | Ile | Ile | Gly | Arg 190 | Ile | Ser |
| Ala | Glu | Gly 195 | Gly | Glu | Gln | Val | Glu 200 | Arg | Val | Val | Glu | Gly 205 | Pro | Ala | Gln |

Leu Leu Leu Leu Asp Val Val Ser Leu Ala Tyr Glu Ser Lys His Leu 225 230 235 240

Leu Glu Gly Ala Lys Ser Glu Ser Ala Glu Glu Leu Lys Lys Arg Ala 245 250 255

Gln Glu Leu Glu Gly Lys Leu Asn Phe Leu Thr Lys Ile His Glu Met 260 265 270

Leu Gln Pro Gly Gln Asp Gln 275

<210> 242

<211> 414

<212> PRT

<213> Homo sapiens

<400> 242

Met Arg Phe Lys Ser His Thr Val Glu Leu Arg Arg Pro Cys Ser Asp 1 5 10 15

Met Glu Gly Ala Ala Leu Leu Arg Val Ser Val Leu Cys Ile Trp Met
20 25 30

Ser Ala Leu Phe Leu Gly Val Arg Val Arg Ala Glu Glu Ala Gly Ala 35 · 40 45

Arg Val Gln Gln Asn Val Pro Ser Gly Thr Asp Thr Gly Asp Pro Gln 50 55 60

Ser Lys Pro Leu Gly Asp Trp Ala Ala Gly Thr Met Asp Pro Glu Ser 65 70 75 80

Ser Ile Phe Ile Glu Asp Ala Ile Lys Tyr Phe Lys Glu Lys Val Ser 85 90 95

Thr Gln Asn Leu Leu Leu Leu Thr Asp Asn Glu Ala Trp Asn Gly
100 105 110

Phe Val Ala Ala Ala Glu Leu Pro Arg Asn Glu Ala Asp Glu Leu Arg 115 120 125

Lys Ala Leu Asp Asn Leu Ala Arg Gln Met Ile Met Lys Asp Lys Asn 130 135 140

Trp His Asp Lys Gly Gln Gln Tyr Arg Asn Trp Phe Leu Lys Glu Phe

| 150 | 155 | 160 |
|-----|-----|-----|
| | | |

| Pro Arg Leu Lys | Ser Lys Leu | Glu Asp Asn Ile | Arg Arg Leu Arg Ala |
|-----------------|-------------|-----------------|---------------------|
| | 165 | 170 | 175 |

- Leu Ala Asp Gly Val Gln Lys Val His Lys Gly Thr Thr Ile Ala Asn 180 185 190
- Val Val Ser Gly Ser Leu Ser Ile Ser Ser Gly Ile Leu Thr Leu Val 195 200 205
- Gly Met Gly Leu Ala Pro Phe Thr Glu Gly Gly Ser Leu Val Leu Leu 210 215 220
- Glu Pro Gly Met Glu Leu Gly Ile Thr Ala Ala Leu Thr Gly Ile Thr 225 230 235 240
- Ser Ser Thr Ile Asp Tyr Gly Lys Lys Trp Trp Thr Gln Ala Gln Ala 245 250 255
- His Asp Leu Val Ile Lys Ser Leu Asp Lys Leu Lys Glu Val Lys Glu 260 265 270
- Phe Leu Gly Glu Asn Ile Ser Asn Phe Leu Ser Leu Ala Gly Asn Thr 275 280 285
- Tyr Gln Leu Thr Arg Gly Ile Gly Lys Asp Ile Arg Ala Leu Arg Arg 290 295 300
- Ala Arg Ala Asn Leu Gln Ser Val Pro His Ala Ser Ala Ser Arg Pro 305 310 315 320
- Arg Val Thr Glu Pro Ile Ser Ala Glu Ser Gly Glu Gln Val Glu Arg 325 330 335
- Val Asn Glu Pro Ser Ile Leu Glu Met Ser Arg Gly Val Lys Leu Thr 340 345 350
- Asp Val Ala Pro Val Ser Phe Phe Leu Val Leu Asp Val Val Tyr Leu 355 360 365
- Val Tyr Glu Ser Lys His Leu His Glu Gly Ala Lys Ser Glu Thr Ala 370 375 380
- Glu Glu Leu Lys Lys Val Ala Gln Glu Leu Glu Glu Lys Leu Asn Ile 385 390 395 400
- Leu Asn Asn Tyr Lys Ile Leu Gln Ala Asp Gln Glu Leu

<210> 243

<211> 398

<212> PRT

<213> Homo sapiens

<400> 243

Met Glu Gly Ala Ala Leu Leu Arg Val Ser Val Leu Cys Ile Trp Met

1 5 10 15

Ser Ala Leu Phe Leu Gly Val Gly Val Arg Ala Glu Glu Ala Gly Ala 20 25 30

Arg Val Gln Gln Asn Val Pro Ser Gly Thr Asp Thr Gly Asp Pro Gln 35 40 45

Ser Lys Pro Leu Gly Asp Trp Ala Ala Gly Thr Met Asp Pro Glu Ser 50 55 60

Ser Ile Phe Ile Glu Asp Ala Ile Lys Tyr Phe Lys Glu Lys Val Ser 65 70 75 80

Thr Gln Asn Leu Leu Leu Leu Thr Asp Asn Glu Ala Trp Asn Gly
85 90 95

Phe Val Ala Ala Ala Glu Leu Pro Arg Asn Glu Ala Asp Glu Leu Arg 100 105 110

Lys Ala Leu Asp Asn Leu Ala Arg Gln Met Ile Met Lys Asp Lys Asn 115 120 125

Trp His Asp Lys Gly Gln Gln Tyr Arg Asn Trp Phe Leu Lys Glu Phe 130 135 140

Pro Arg Leu Lys Ser Glu Leu Glu Asp Asn Ile Arg Arg Leu Arg Ala 145 150 155 160

Leu Ala Asp Gly Val Gln Lys Val His Lys Gly Thr Thr Ile Ala Asn 165 170 175

Val Val Ser Gly Ser Leu Ser Ile Ser Ser Gly Ile Leu Thr Leu Val 180 185 190

Gly Met Gly Leu Ala Pro Phe Thr Glu Gly Gly Ser Leu Val Leu Leu 195 200 205

Glu Pro Gly Met Glu Leu Gly Ile Thr Ala Ala Leu Thr Gly Ile Thr Ser Ser Thr Ile Asp Tyr Gly Lys Lys Trp Trp Thr Gln Ala Gln Ala His Asp Leu Val Ile Lys Ser Leu Asp Lys Leu Lys Glu Val Lys Glu Phe Leu Gly Glu Asn Ile Ser Asn Phe Leu Ser Leu Ala Gly Asn Thr Tyr Gln Leu Thr Arg Gly Ile Gly Lys Asp Ile Arg Ala Leu Arg Arg Ala Arg Ala Asn Leu Gln Ser Val Pro His Ala Ser Ala Ser Arg Pro Arg Val Thr Glu Pro Ile Ser Ala Glu Ser Gly Glu Gln Val Glu Arg Val Asn Glu Pro Ser Ile Leu Glu Met Ser Arg Gly Val Lys Leu Thr Asp Val Ala Pro Val Gly Phe Phe Leu Val Leu Asp Val Val Tyr Leu Val Tyr Glu Ser Lys His Leu His Glu Gly Ala Lys Ser Glu Thr Ala Glu Glu Leu Lys Lys Val Ala Gln Glu Leu Glu Glu Lys Leu Asn Met Leu Asn Asn Asn Tyr Lys Ile Leu Gln Ala Asp Gln Glu Leu <210> 244 <211> 479 <212> PRT <213> Homo sapiens <400> 244 Met Ala Trp Asn Thr Asn Leu Arg Trp Arg Leu Pro Leu Thr Cys Leu Leu Leu Gln Val Ile Met Val Ile Leu Phe Gly Val Phe Val Arg Tyr

Asp Phe Glu Ala Asp Ala His Trp Trp Ser Glu Arg Thr His Lys Asn Leu Ser Asp Met Glu Asn Glu Phe Tyr Tyr Arg Tyr Pro Ser Phe Gln Asp Val His Val Met Val Phe Val Gly Phe Gly Phe Leu Met Thr Phe Leu Gln Arg Tyr Gly Phe Ser Ala Val Gly Phe Asn Phe Leu Leu Ala Ala Phe Gly Ile Gln Trp Ala Leu Leu Met Gln Gly Trp Phe His Phe Leu Gln Asp Arg Tyr Ile Val Val Gly Val Glu Asn Leu Ile Asn Ala Asp Phe Cys Val Ala Ser Val Cys Val Ala Phe Gly Ala Val Leu Gly Lys Val Ser Pro Ile Gln Leu Leu Ile Met Thr Phe Phe Gln Val Thr Leu Phe Ala Val Asn Glu Phe Ile Leu Leu Asn Leu Leu Lys Val Lys Asp Ala Gly Gly Ser Met Thr Ile His Thr Phe Gly Ala Tyr Phe Gly Leu Thr Val Thr Arg Ile Leu Tyr Arg Arg Asn Leu Glu Gln Ser Lys Glu Arg Gln Asn Ser Val Tyr Gln Ser Asp Leu Phe Ala Met Ile Gly Thr Leu Phe Leu Trp Met Tyr Trp Pro Ser Phe Asn Ser Ala Ile Ser Tyr His Gly Asp Ser Gln His Arg Ala Ala Ile Asn Thr Tyr Cys Ser Leu Ala Ala Cys Val Leu Thr Ser Val Ala Ile Ser Ser Ala Leu His Lys Lys Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu Ala

Gly Gly Val Ala Val Gly Thr Ala Ala Glu Met Met Leu Met Pro Tyr Gly Ala Leu Ile Ile Gly Phe Val Cys Gly Ile Ile Ser Thr Leu Gly Phe Val Tyr Leu Thr Pro Phe Leu Glu Ser Arg Leu His Ile Gln Asp Thr Cys Gly Ile Asn Asn Leu His Gly Ile Pro Gly Ile Ile Gly Gly Ile Val Gly Ala Val Thr Ala Ala Ser Ala Ser Leu Glu Val Tyr Gly Lys Glu Gly Leu Val His Ser Phe Asp Phe Gln Gly Phe Asn Gly Asp Trp Thr Ala Arg Thr Gln Gly Lys Phe Gln Ile Tyr Gly Leu Leu Val Thr Leu Ala Met Ala Leu Met Gly Gly Ile Ile Val Gly Leu Ile Leu Arg Leu Pro Phe Trp Gly Gln Pro Ser Asp Glu Asn Cys Phe Glu Asp Ala Val Tyr Trp Glu Met Pro Glu Gly Asn Ser Thr Val Tyr Ile Pro Glu Asp Pro Thr Phe Lys Pro Ser Gly Pro Ser Val Pro Ser Val Pro Met Val Ser Pro Leu Pro Met Ala Ser Ser Val Pro Leu Val Pro <210> 245 <211> 498 <212> PRT <213> Mus musculus <400> 245 Met Ala Trp Asn Thr Asn Leu Arg Gly Arg Leu Pro Ile Thr Cys Leu

Ile Leu Gln Val Thr Met Val Val Leu Phe Gly Val Phe Val Arg Tyr

| 20 | 25 | 30 |
|----|----|----|
| | | |

Asp Ile Gln Ala Asp Ala His Trp Trp Leu Glu Lys Lys Arg Lys Asn

- Ile Ser Ser Asp Val Glu Asn Glu Phe Tyr Tyr Arg Tyr Pro Ser Phe
- Gln Asp Val His Ala Met Val Phe Val Gly Phe Gly Phe Leu Met Thr .
- Phe Leu Gln Arg Tyr Gly Phe Ser Ala Val Gly Phe Asn Phe Leu Leu
- Ala Ala Phe Gly Ile Gln Trp Ala Leu Leu Met Gln Gly Trp Phe His
- Tyr Phe Glu Glu Gly His Ile Val Leu Ser Val Glu Asn Ile Ile Gln
- Ala Asp Phe Cys Val Ala Ser Ser Cys Val Ala Phe Gly Ala Val Leu
- Gly Lys Val Ser Pro Met Gln Leu Leu Ile Met Thr Phe Phe Gln Val
- Thr Leu Phe Thr Val Asn Glu Phe Ile Leu Leu Asn Leu Ile Glu Ala
- Lys Asp Ala Gly Gly Ser Met Thr Ile His Thr Phe Gly Ala Tyr Phe
- Gly Leu Thr Val Thr Trp Ile Leu Tyr Arg Lys Asn Leu Asp Gln Ser
- Lys Gln Arg Gln Ser Ser Val Tyr His Ser Asp Leu Phe Ala Met Ile
- Gly Thr Leu Phe Leu Trp Ile Tyr Trp Pro Ser Phe Asn Ser Ala Ser
- Ser Phe His Gly Asp Ala Gln His Arg Ala Ala Leu Asn Thr Tyr Leu
- Ser Leu Ala Ala Ser Val Leu Thr Thr Val Thr Val Ser Ser Ile Val
- His Lys Lys Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu

- Ala Gly Gly Val Gly Val Gly Thr Ala Ala Glu Met Met Leu Thr Pro 290 295 300
- Tyr Gly Ala Leu Ile Val Gly Phe Phe Cys Gly Ile Phe Ser Thr Leu 305 310 315 320
- Gly Phe Ala Tyr Leu Thr Pro Phe Leu Glu Ser Arg Leu Arg Ile Gln 325 330 335
- Asp Thr Cys Gly Ile His Asn Leu His Gly Ile Pro Gly Ile Ile Gly 340 345 350
- Gly Ile Val Gly Ala Val Thr Ala Ala Tyr Ser Ser Pro Asp Val Tyr 355 360 365
- Gly Glu Pro Gly Ile Val His Ser Phe Gly Phe Gly Ser Tyr Lys Met 370 380
- Asp Trp Asn Lys Arg Met Gln Gly Arg Ser Gln Ile Phe Gly Leu Leu 385 390 395 400
- Leu Ser Leu Ala Met Ala Leu Val Gly Gly Ile Ile Val Gly Phe Ile 405 410 415
- Leu Lys Leu Pro Phe Trp Gly Gln Ala Ala Asp Glu Asn Cys Phe Glu
 420 425 430
- Asp Ser Ile Tyr Trp Glu Val His Glu Glu Val Asn Thr Val Tyr Ile
 435 440 445
- Pro Glu Asp Leu Ala His Lys His Ser Thr Ser Leu Val Pro Ala Met 450 455 460
- Pro Leu Val Leu Pro Thr Thr Ser Ala Ser Ile Val Pro Pro Val Pro 465 470 475 480
- Pro Thr Pro Pro Val Ser Leu Ala Thr Ser Ala Pro Ser Ala Ala Leu 485 490 495

Val His

<210> 246

<211> 459

<212> PRT

<213> Bos taurus

210

225

| <400 |)> 24 | 16 | | | | | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Met 1 | Ile | Trp | Asn | Thr 5 | Asn | Leu | Arg | Trp | Arg 10 | Leu | Pro | Val | Ala | Cys 15 | Leu |
| Leu | Leu | Glu | Val 20 | Ala | Leu | Ile | Ala | Leu 25 | Phe | Gly | Val | Phe | Val 30 | Arg | Tyr |
| Asp | Met | Asp 35 | Ala | Asp | Pro | His | Trp | Val | Gln | Glu | Lys | Val 45 | Ile | Lys | Asn |
| Leu | Ser 50 | Thr | Asp | Leu | Glu | Asn 55 | Glu | Phe | Tyr | Tyr | Arg 60 | Tyr | Pro | Ser | Phe |
| Gln 65 | Asp | Val | His | Val | Met 70 | Ile | Phe | Val | Gly | Phe 75 | Gly | Phe | Leu | Met | Thr 80 |
| Phe | Leu | Gln | Arg | Tyr 85 | Gly | Tyr | Ser | Ser | Val 90 | Gly | Phe | Asn | Phe | Leu 95 | Ala |
| Ala | Phe | Gly | Ile 100 | Gln | Trp | Ala | Leu | Leu 105 | Met | Gln | Gly | Trp | Leu 110 | Gln | Ser |
| Phe | Asp | Gly 115 | Arg | Tyr | Ile | Leu | Val 120 | Asp | Leu | Glu | Asn | Leu 125 | Ile | Asn | Ala |
| Asp | Phe 130 | Cys | Val | Gly | Ser | Val 135 | Cys | Val | Ala | Phe | Gly 140 | Ala | Val | Leu | Gly |
| Lys 145 | Val | Ser | Pro | Val | Gln 150 | Leu | Leu | Ile | Met | Thr 155 | Leu | Phe | Gln | Val | Thr 160 |
| Leu | Phe | Ser | Ile | Asn 165 | Glu | Tyr | Ile | Leu | Leu 170 | Asn | Leu | Leu | Glu | Val 175 | Lys |
| Asp | Ser | Gly | Gly 180 | Ser | Met | Thr | Ile | His 185 | Ala | Phe | Gly | Ala | Tyr 190 | Phe | Gly |
| Leu | Thr | Val 195 | Ala | Trp | Ile | Leu | Tyr 200 | Arg | Pro | Asn | Leu | His 205 | Leu | Ser | Lys |
| Glu | Arg | Gln | Ser | Ser | Thr | Tyr | His | Ser | Asp | Leu | Phe | Ala | Met | Ile | Gly |

Thr Leu Phe Leu Trp Met Tyr Trp Pro Ser Phe Asn Ser Ala Ile Ser

235 240

215 220

Asn His Gly Asp Ala Gln His Arg Ala Ile Asn Thr Tyr Cys Ser Leu Ala Ala Cys Val Leu Thr Ser Val Ala Leu Ser Ser Ala Leu His Arg Lys Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu Ala Gly Gly Val Gly Leu Gly Thr Val Ala Glu Leu Met Val Leu Pro Phe Gly Ser Leu Ile Ile Gly Phe Val Cys Gly Ile Val Ser Thr Leu Gly Phe Val Tyr Leu Thr Pro Phe Leu Glu Ser Arg Leu His Ile Gln Asp Thr Cys Gly Val His Asn Leu His Gly Ile Pro Gly Ile Ile Gly Gly Ile Ala Gly Ala Val Thr Ala Ser Ile Ala Asn Ile Asp Leu Tyr Gly Glu Glu Gly Leu Ala Tyr Ala Phe Gly Ile Glu Arg Ser Lys Leu Asn Trp Ser Pro Asn Met Gln Gly Arg Phe Gln Ala Ala Gly Leu Phe Val Ser Leu Ala Met Ala Leu Val Gly Gly Val Ile Val Gly Val Ile Leu Arg Leu Pro Phe Trp Gly Gln Ala Pro Asp Glu Asn Cys Phe Glu Asp

Ala Val Tyr Trp Glu Ile Pro Lys Glu Pro Lys Ser Thr Ala Leu Arg 435 440 445

Ser Glu Asp Ser Ser Ile Lys Pro Pro Glu Pro 450 455

<210> 247

<211> 467

<212> PRT

<213> Orycctolagus cuniculus

| <4 | 00> | 247 |
|----|-----|-----|
| | | |

- Met Ala Trp Asn Thr Asn Leu Arg Trp Arg Leu Pro Leu Leu Cys Leu 1 5 10 15
- Val Leu Glu Val Ala Met Val Val Leu Phe Gly Leu Phe Val Arg Tyr
 20 25 30
- Ser Pro Asp Ala Asp Ser Ser Trp Ser Asn Glu Lys Arg Lys Gly Asn 35 40 45
- Ile Thr Ser Asp Leu Glu Asn Glu Phe Tyr Tyr Arg Tyr Pro Ser Phe 50 55 60
- Gln Asp Val His Val Met Val Phe Leu Gly Phe Gly Phe Leu Met Thr 65 70 75 80
- Phe Leu Gln Arg Tyr Gly Tyr Cys Ala Leu Gly Phe Asn Phe Leu Leu 85 90 95
- Ala Ala Leu Gly Val Gln Trp Ala Leu Leu Met Gln Gly Trp Phe Gln
 100 105 110
- Tyr Thr Lys Asp Arg Leu Ile Leu Leu Gly Ile Lys Asn Leu Ile Asp 115 120 125
- Ala Asp Ser Cys Val Ala Ser Val Cys Val Ala Phe Gly Ala Val Leu 130 135 140
- Gly Lys Val Ser Pro Val Gln Met Leu Leu Met Thr Phe Phe Gln Val 145 150 155 160
- Ala Leu Phe Ser Ala Asn Glu Phe Leu Leu His Val Leu Glu Val 165 170 175
- Lys Asp Ala Gly Gly Ser Ile Thr Ile His Ile Phe Gly Ala Tyr Phe 180 185 190
- Gly Leu Thr Val Thr Trp Ile Leu Tyr Arg His Asn Leu Asp His Ser 195 200 205
- Arg Glu Arg Gln Ser Ser Val Tyr His Ser Asn Leu Phe Ala Met Ile 210 215 220
- Gly Thr Leu Phe Leu Trp Ile Tyr Trp Pro Ser Phe Asn Ser Ala Met 225 230 235 240
- Ser Asn Tyr Gly Asp Ala Gln His Arg Ala Ala Ile Asn Thr Tyr Cys 245 250 255

Ser Leu Ala Ala Ser Val Leu Thr Ser Val Ala Met Ser Ser Val Leu 270

His Lys Lys Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu 280

Ala Gly Gly Val Gly Val Gly Thr Ala Ala Glu Glu Met 300

Tyr Gly Ala Leu Ile Val Gly Phe Ile Cys Gly Ala Val Ser Thr Leu 305

Gly Phe Val Tyr Leu Thr Pro Phe Leu Glu Ser Arg Leu Arg Ile Gln 325 330 335

Asp Thr Cys Gly Ile His Asn Leu His Gly Ile Pro Gly Leu Ile Gly 340 345 350

Ala Ile Val Gly Ala Val Thr Ala Ala Tyr Ala Ser Pro Asp Gly Asp 355 360 365

Arg Gly Phe Val Tyr Pro Phe Gly Phe His Asn Glu Lys Asp Glu Lys 370 375 380

Val Gln Gly Arg Phe Gln Ala Phe Gly Leu Leu Thr Leu Ala Ile 385 390 395 400

Ala Met Val Gly Gly Thr Ile Met Gly Leu Ile Leu Lys Leu Pro Phe 405 410 415

Trp Gly Gln Ala Met Asp Glu Asp Cys Phe Asp Asp Ser Ile Tyr Trp
420 425 430

Glu Met His Glu Glu Lys Ser Ser Pro Glu Asp His Thr His Lys 435 440 445

Pro Ser Val Pro Thr Glu Pro Val Glu Gln Pro Thr Ser Ser Ala Thr 450 455 460

Leu Ala Pro 465

<210> 248

<211> 488

<212> PRT

<213> Oryzias latipes

- Met Gly Asn Cys Cys Glu Ser Ala Ser Asn Phe Phe Gly Pro Gln Lys
 1 5 10 15
- Asn Thr Asn Val Arg Val Ser Leu Pro Ala Val Cys Phe Val Trp Gln
 20 25 30
- Ile Ala Met Ile Val Leu Phe Gly Val Phe Ile Arg Tyr Asp Glu Glu
 35 40 45
- Ser Asp Ala His Trp Val Glu Leu Lys Lys Thr Glu Asn Leu Thr Asp 50 55 60
- Leu Gln Asn Glu Phe Tyr Phe Arg Tyr Pro Ser Phe Gln Asp Val His
 65 70 75 80
- Val Met Ile Phe Val Gly Phe Gly Phe Leu Met Thr Phe Leu Lys Arg 85 90 95
- Tyr Ser Phe Ser Ala Val Gly Phe Asn Phe Leu Ile Ala Ala Phe Gly
 100 105 110
- Leu Gln Trp Ala Leu Leu Met Gln Gly Trp Phe His His Phe Asp Tyr
 115 120 125
- Ser Thr Gly Lys Ile Tyr Ile Gly Ile Glu Ser Leu Ile Asn Ala Asp 130 135 140
- Phe Cys Cys Ala Ala Ser Leu Ile Ala Tyr Gly Ala Ile Leu Gly Lys 145 150 155 160
- Val Ser Pro Val Gln Leu Met Val Val Thr Leu Phe Gly Val Thr Leu
 165 170 175
- Phe Ala Val Glu Glu Tyr Ile Ile Leu Asp Leu Leu His Cys Arg Asp 180 185 190
- Ser Gly Gly Ala Met Val Ile His Cys Phe Gly Gly Tyr Tyr Gly Leu 195 200 205
- Ala Ile Ser Trp Val Leu Tyr Arg Pro Asn Leu His Arg Ser Lys Arg 210 215 220
- Leu Asn Gly Ser Val Tyr His Ser Asp Leu Phe Ala Met Ile Gly Thr 225 230 235 240
- Leu Phe Leu Trp Met Phe Trp Pro Ser Phe Asn Ser Ala Ile Ala Asn

- His Gly Asp Gly Gln His Arg Thr Ala Met Asn Thr Tyr Ile Ala Leu 260 265 270
- Ala Ser Ser Val Leu Thr Thr Val Ala Leu Ser Ser Met Ser Lys Lys 275 280 285
- Glu Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu Ala Gly 290 295 300
- Gly Val Ala Met Gly Thr Ala Ala Glu Phe Met Ile Thr Pro Tyr Gly 315 310 315
- Ser Leu Ile Val Gly Phe Cys Ile Gly Ile Ile Ser Thr Phe Gly Tyr 325 330 335
- Leu Tyr Val Thr Pro Phe Leu Glu Lys Arg Leu Lys Leu Gln Asp Thr 340 345 350
- Cys Gly Ile His Asn Leu His Ala Val Pro Gly Met Leu Gly Gly Phe 355 360 365
- Ile Gly Ala Ile Val Ala Ala Thr Ala Ser Glu Ser Val Tyr Ser Lys 370 375 380
- Gln Gly Leu Ile Asp Thr Phe Gly Phe Thr Gly Lys Tyr Glu Asn Arg 385 390 395 400
- Ser Pro Gly Thr Gln Gly Gly Tyr Gln Ala Ala Gly Val Cys Val Ala 405 410 415
- Met Ala Phe Gly Leu Val Gly Gly Ala Ile Val Gly Phe Ile Leu Lys 420 425 430
- Phe Pro Ile Trp Gly Asp Ala Ala Asp Asp Tyr Cys Phe Asp Asp Glu 435 440 445
- Ala Tyr Trp Glu Leu Pro Glu Glu Glu Glu Thr Ile Pro Pro Val Leu 450 455 460
- Glu Tyr Asn Asn His Met Thr Gln Gln Lys His Gln Glu Thr Pro Glu 465 470 475 480

Thr Ser Phe Ser Val Val Glu Ser 485

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<210> 249
<211> 388
<212> PRT
<213> Homo sapiens
         35
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- Ala Ala Phe Gly Ile Gln Trp Ala Leu Leu Met Gln Gly Trp Phe His
- Phe Leu Gln Asp Arg Tyr Ile Val Val Gly Val Glu Asn Leu Ile Asn
- Ala Asp Phe Cys Val Ala Ser Val Cys Val Ala Phe Gly Ala Val Leu
- Gly Lys Val Ser Pro Ile Gln Leu Leu Ile Met Thr Phe Phe Gln Val
- Thr Leu Phe Ala Val Asn Glu Phe Ile Leu Leu Asn Leu Leu Lys Val
- Lys Asp Ala Gly Gly Ser Met Thr Ile His Thr Phe Gly Ala Tyr Phe
- Gly Leu Thr Val Thr Arg Ile Leu Tyr Arg Arg Asn Leu Glu Gln Ser
- Lys Glu Arg Gln Asn Ser Val Tyr Gln Ser Asp Leu Phe Ala Met Ile
- Gly Thr Leu Phe Leu Trp Met Tyr Trp Pro Ser Phe Asn Ser Ala Ile
- Ser Tyr His Gly Asp Ser Gln His Arg Ala Ile Asn Thr Tyr Cys
- Ser Leu Ala Ala Cys Val Leu Thr Ser Val Ala Ile Ser Ser Ala Leu

His Lys Lys Gly Lys Leu Asp Met Val His Ile Gln Asn Ala Thr Leu Ala Gly Gly Val Ala Val Gly Thr Ala Ala Glu Met Met Leu Met Pro Tyr Gly Ala Leu Ile Ile Gly Phe Val Cys Gly Ile Ile Ser Thr Leu Gly Phe Val Tyr Leu Thr Pro Phe Leu Glu Ser Arg Leu His Ile Gln Asp Thr Cys Gly Ile Asn Asn Leu His Gly Ile Pro Gly Ile Ile Gly Gly Ile Val Gly Ala Val Thr Ala Ala Ser Ala Ser Leu Glu Val Tyr Gly Lys Glu Gly Leu Val His Ser Phe Asp Phe Gln Gly Phe Asn Gly Asp Trp Thr Ala Arg Thr Gln Gly Lys Phe Gln Ile Tyr Gly Leu Leu Val Thr Leu Ala Met Ala Leu Met Gly Gly Ile Ile Val Gly Leu Ile Leu Arg Leu Pro Phe Trp Gly Gln Pro Ser Asp Glu Asn Cys Phe Glu Asp Ala Val Tyr <210> 250 <211> 373 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Ammonium Transporter Family domain sequence <400> 250 Gly Leu Val Arg Ser Lys Asn Val Leu Asn Ile Leu Tyr Lys Asn Phe Gln Asp Val Ala Ile Gly Val Leu Ala Tyr Trp Gly Phe Gly Tyr Ser

30

- Leu Ala Phe Gly Asp Ser Tyr Phe Ser Gly Phe Ile Gly Asn Leu Gly 35 40 45
- Leu Leu Ala Ala Gly Ile Gln Trp Gly Thr Leu Pro Asp Gly Leu Phe 50 55 60
- Phe Leu Phe Gln Leu Met Phe Ala Ala Thr Ala Ile Thr Ile Ile Ser 65 70 75 80
- Gly Ala Val Ala Glu Arg Ile Lys Phe Ser Ala Tyr Leu Leu Phe Ser 85 90 95
- Ala Leu Leu Gly Thr Leu Val Tyr Pro Pro Val Ala His Trp Val Trp
 100 105 110
- Gly Glu Gly Gly Trp Leu Ala Lys Leu Gly Val Leu Val Asp Phe Ala 115 120 125
- Gly Ser Thr Val Val His Ile Phe Gly Gly Tyr Ala Gly Leu Ala Ala 130 135 140
- Ala Leu Val Leu Gly Pro Arg Ile Gly Arg Phe Thr Lys Asn Glu Ala 145 150 155 160
- Ile Thr Pro His Asn Leu Pro Phe Ala Val Leu Gly Thr Leu Leu Leu 165 170 175
- Trp Phe Gly Trp Phe Gly Phe Asn Ala Gly Ser Ala Leu Thr Ala Asp 180 185 190
- Gly Arg Ala Arg Ala Ala Ala Val Asn Thr Asn Leu Ala Ala Gly
 195 200 205
- Gly Ala Leu Thr Ala Leu Leu Ile Ser Arg Leu Lys Thr Gly Lys Pro 210 215 220
- Asn Met Leu Gly Leu Ala Asn Gly Ala Leu Ala Gly Leu Val Ala Ile 225 230 235 240
- Thr Pro Ala Cys Gly Val Val Ser Pro Trp Gly Ala Leu Ile Ile Gly 245 250 255
- Leu Ile Ala Gly Val Leu Ser Val Leu Gly Tyr Lys Leu Lys Glu Lys 260 265 270
- Leu Gly Ile Asp Asp Pro Leu Asp Val Phe Pro Val His Gly Val Gly

275 280 285

Gly Ile Trp Gly Gly Ile Ala Val Gly Ile Phe Ala Ala Leu Tyr Val 290 295 300

Asn Thr Ser Gly Ile Tyr Gly Gly Leu Leu Tyr Gly Asn Ser Lys Gln 305 310 315 320

Leu Gly Val Gln Leu Ile Gly Ile Ala Val Ile Leu Ala Tyr Ala Phe 325 330 335

Gly Val Thr Phe Ile Leu Gly Leu Leu Gly Leu Thr Leu Gly Leu 340 345 350

Arg Val Ser Glu Glu Glu Glu Lys Val Gly Leu Asp Leu Ala Glu His 355 360 365

Gly Glu Thr Ala Tyr 370

<210> 251

<211> 446

<212> PRT

<213> Homo sapiens

<400> 251

Met Arg Leu Asp Glu His Asp Phe Leu Gly Gln Phe Ser Cys Ser Leu 1 5 10 15

Gly Thr Ile Val Ser Ser Lys Lys Ile Thr Arg Pro Leu Leu Leu 20 25 30

Asn Asp Lys Pro Ala Gly Lys Gly Leu Ile Thr Ile Ala Ala Gln Glu 35 40 45

Leu Ser Asp Asn Arg Val Ile Thr Leu Ser Leu Ala Gly Arg Arg Leu 50 55 60

Asp Lys Lys Asp Leu Phe Gly Lys Ser Asp Pro Phe Leu Glu Phe Tyr 65 70 75 80

Lys Pro Gly Asp Asp Gly Lys Trp Met Leu Val His Arg Thr Glu Val 85 90 95

Ile Lys Tyr Thr Leu Asp Pro Val Trp Lys Pro Phe Thr Val Pro Leu 100 105 . 110

| Val | Ser | Leu 115 | Cys | Asp | Gly | Asp | Met 120 | Glu | Lys | Pro | Ile | Gln 125 | Val | Met | Cys |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Tyr | Asp 130 | Tyr | Asp | Asn | Asp | Gly 135 | Gly | His | Asp | Phe | Ile 140 | Gly | Glu | Phe | Glr |
| Thr 145 | Ser | Val | Ser | Gln | Met 150 | Cys | Glu | Ala | Arg | Asp 155 | Ser | Val | Pro | Leu | Glu 160 |
| Phe | Glu | Cys | Ile | Asn 165 | Pro | Lys | Lys | Gln | Arg 170 | Lys | Lys | Lys | Asn | Tyr 175 | Lys |
| Asn | Ser | Gly | Ile 180 | Ile | Ile | Leu | Arg | Ser 185 | Cys | Lys | Ile | Asn | Arg 190 | Asp | Tyr |
| Ser | Phe | Leu 195 | Asp | Tyr | Ile | Leu | Gly 200 | Gly | Cys | Gln | Leu | Met 205 | Phe | Thr | Val |
| Gly | Ile 210 | Asp | Phe | Thr | Ala | Ser 215 | Asn | Gly | Asn | Pro | Leu 220 | Asp | Pro | Ser | Ser |
| Leu 225 | His | Tyr | Ile | Asn | Pro 230 | Met | Gly | Thr | Asn | Glu 235 | Tyr | Leu | Ser | Ala | 11e |
| Trp | Ala | Val | Gly | Gln 245 | Ile | Ile | Gln | Asp | Tyr 250 | Asp | Ser | Asp | Lys | Met 255 | Phe |
| Pro | Ala | Leu | Gly 260 | Phe | Gly | Ala | Gln | Leu 265 | Pro | Pro | Asp | Trp | Lys 270 | Val | Ser |
| His | Glu | Phe 275 | Ala | Ile | Asn | Phe | Asn 280 | Pro | Thr | Asn | Pro | Phe 285 | Cys | Ser | Gly |
| Val | Asp 290 | Gly | Ile | Ala | Gln | Ala 295 | Tyr | Ser | Ala | Cys | Leu 300 | Pro | His | Ile | Arg |
| Phe 305 | Tyr | Gly | Pro | Thr | Asn 310 | Phe | Ser | Pro | Ile | Val 315 | Asn | His | Val | Ala | Arg 320 |
| Phe | Ala | Ala | Gln | Ala 325 | Thr | Gln | Gln | Arg | Thr 330 | Ala | Thr | Gln | Tyr | Phe 335 | Ile |
| Leu | Leu | Ile | Ile 340 | Thr | Asp | Gly | Val | Ile 345 | Ser | Asp | Met | Glu | Glu 350 | Thr | Arg |
| His | Ala | Val | Val | Gln | Ala | Ser | Lys | Leu | Pro | Met | Ser | Ile | Ile | Ile | Val |

Gly Val Gly Asn Ala Asp Phe Ala Ala Met Glu Phe Leu Asp Gly Asp Ser Arg Met Leu Arg Ser His Thr Gly Glu Glu Ala Ala Arg Asp Ile Val Gln Phe Val Pro Phe Arg Glu Phe Arg Asn Ala Ala Lys Glu Thr Leu Ala Lys Ala Val Leu Ala Glu Leu Pro Gln Gln Val Val Gln Tyr Phe Lys His Lys Asn Leu Pro Pro Thr Asn Ser Glu Pro Ala <210> 252 <211> 358 <212> PRT <213> Homo sapiens <400> 252 Met Leu Val His Arg Thr Glu Val Ile Lys Tyr Thr Leu Asp Pro Val Trp Lys Pro Phe Thr Val Pro Leu Val Ser Leu Cys Asp Gly Asp Met Glu Lys Pro Ile Gln Val Met Cys Tyr Asp Tyr Asp Asn Asp Gly Gly His Asp Phe Ile Gly Glu Phe Gln Thr Ser Val Ser Gln Met Cys Glu Ala Arg Asp Ser Val Pro Leu Glu Phe Glu Cys Ile Asn Pro Lys Lys Gln Arg Lys Lys Asn Tyr Lys Asn Ser Gly Ile Ile Leu Arg Ser Cys Lys Ile Asn Arg Asp Tyr Ser Phe Leu Asp Tyr Ile Leu Gly Gly Cys Gln Leu Met Phe Thr Val Gly Ile Asp Phe Thr Ala Ser Asn Gly Asn Pro Leu Asp Pro Ser Ser Leu His Tyr Ile Asn Pro Met Gly

Asp Tyr Asp Ser Asp Lys Met Phe Pro Ala Leu Gly Phe Gly Ala Gln 165 170 175

Leu Pro Pro Asp Trp Lys Val Ser His Glu Phe Ala Ile Asn Phe Asn 180 185 190

Pro Thr Asn Pro Phe Cys Ser Gly Val Asp Gly Ile Ala Gln Ala Tyr 195 200 205

Ser Ala Cys Leu Pro His Ile Arg Phe Tyr Gly Pro Thr Asn Phe Ser 210 215 220

Pro Ile Val Asn His Val Ala Arg Phe Ala Ala Gln Ala Thr Gln Gln 225 230 235 240

Arg Thr Ala Thr Gln Tyr Phe Ile Leu Leu Ile Ile Thr Asp Gly Val 245 250 255

Ile Ser Asp Met Glu Glu Thr Arg His Ala Val Val Gln Ala Ser Lys 260 265 270

Leu Pro Met Ser Ile Ile Ile Val Gly Val Gly Asn Ala Asp Phe Ala 275 280 285

Ala Met Glu Phe Leu Asp Gly Asp Ser Arg Met Leu Arg Ser His Thr 290 295 300

Gly Glu Glu Ala Ala Arg Asp Ile Val Gln Phe Val Pro Phe Arg Glu 305 310 315 320

Phe Arg Asn Ala Ala Lys Glu Thr Leu Ala Lys Ala Val Leu Ala Glu 325 330 335

Leu Pro Gln Gln Val Val Gln Tyr Phe Lys His Lys Asn Leu Pro Pro 340 345 350

Thr Asn Ser Glu Pro Ala 355

<210> 253

<211> 537

<212> PRT

<213> Homo sapiens

| <400> 253 | | | | | | | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Met 1 | Ala | Ala | Gln | Cys 5 | Val | Thr | Lys | Val | Ala 10 | Leu | Asn | Val | Ser | Cys 15 | Ala |
| Asn | Leu | Leu | Asp 20 | Lys | Asp | Ile | Gly | Ser 25 | Lys | Ser | Asp | Pro | Leu 30 | Cys | Val |
| Leu | Phe | Leu 35 | Asn | Thr | Ser | Gly | Gln 40 | Gln | Trp | Tyr | Glu | Val 45 | Glu | Arg | Thr |
| Glu | Arg 50 | Ile | Lys | Asn | Cys | Leu 55 | Asn | Pro | Gln | Phe | Ser 60 | Lys | Thr | Phe | Ile |
| Ile 65 | Asp | Tyr | Tyr | Phe | Glu 70 | Val | Val | Gln | Lys | Leu 75 | Lys | Phe | Gly | Val | Tyr 80 |
| Asp | Ile | Asp | Asn | Lys 85 | Thr | Ile | Glu | Leu | Ser 90 | Asp | Asp | Asp | Phe | Leu 95 | Gly |
| Glu | Cys | Glu | Cys 100 | Thr | Leu | Gly | Gln | Ile 105 | Val | Ser | Ser | Lys | Lys 110 | Leu | Thr |
| Arg | Pro | Leu 115 | Val | Met | Lys | Thr | Gly 120 | Arg | Pro | Ala | Gly | Lys 125 | Gly | Ser | Ile |
| Thr | Ile 130 | Ser | Ala | Glu | Glu | Ile 135 | Lys | Asp | Asn | Arg | Val 140 | Val | Leu | Phe | Glu |
| Met 145 | Glu | Ala | Arg | Lys | Leu 150 | Asp | Asn | Lys | Asp | Leu 155 | Phe | Gly | Lys | Ser | Asp 160 |
| Pro | Tyr | Leu | Glu | Phe 165 | His | Lys | Gln | Thr | Ser 170 | Asp | Gly | Asn | Trp | Leu 175 | Met |
| Val | His | Arg | Thr 180 | Glu | Val | Val | Lys | Asn 185 | Asn | Leu | Asn | Pro | Val 190 | Trp | Arg |
| Pro | Phe | Lys 195 | Ile | Ser | Leu | Asn | Ser 200 | Leu | Cys | Tyr | Gly | Asp 205 | Met | Asp | Lys |
| Thr | Ile 210 | Lys | Val | Glu | Cys | Tyr 215 | Asp | Tyr | Asp | Asn | Asp 220 | Gly | Ser | His | Asp |
| Leu | Ile | Gly | Thr | Phe | Gln | Thr | Thr | Met | Thr | Lys | Leu | Lys | Glu | Ala | Ser |

Arg Ser Ser Pro Val Glu Phe Glu Cys Ile Asn Glu Lys Lys Arg Gln

| 245 | 250 | 255 |
|-----|-----|-----|
| | | |

- Lys Lys Ser Tyr Lys Asn Ser Gly Val Ile Ser Val Lys Gln Cys 260 265 270
- Glu Ile Thr Val Glu Cys Thr Phe Leu Asp Tyr Ile Met Gly Gly Cys 275 280 285
- Gln Leu Asn Phe Thr Val Gly Val Asp Phe Thr Gly Ser Asn Gly Asp 290 295 300
- Pro Arg Ser Pro Asp Ser Leu His Tyr Ile Ser Pro Asn Gly Val Asn 305 310 315 320
- Glu Tyr Leu Thr Ala Leu Trp Ser Val Gly Leu Val Ile Gln Asp Tyr 325 330 335
- Asp Ala Asp Lys Met Phe Pro Ala Phe Gly Phe Gly Ala Gln Ile Pro 340 345 350
- Pro Gln Trp Gln Val Ser His Glu Phe Pro Met Asn Phe Asn Pro Ser 355 360 365
- Asn Pro Tyr Cys Asn Gly Ile Gln Gly Ile Val Glu Ala Tyr Arg Ser 370 375 380
- Cys Leu Pro Gln Ile Lys Leu Tyr Gly Pro Thr Asn Phe Ser Pro Ile 385 390 395 400
- Ile Asn His Val Ala Arg Phe Ala Ala Ala Ala Thr Gln Gln Gln Thr
 405 410 415
- Ala Ser Gln Tyr Phe Val Leu Leu Ile Ile Thr Asp Gly Val Ile Thr
 420 425 430
- Asp Leu Asp Glu Thr Arg Gln Ala Ile Val Asn Ala Ser Arg Leu Pro 435 440 445
- Met Ser Ile Ile Ile Val Gly Val Gly Gly Ala Asp Phe Ser Ala Met 450 455 460
- Glu Phe Leu Asp Gly Asp Gly Gly Ser Leu Arg Ser Pro Leu Gly Glu 465 470 475 480
- Val Ala Ile Arg Asp Ile Val Gln Phe Val Pro Phe Arg Gln Phe Gln 485 490 495
- Asn Ala Pro Lys Glu Ala Leu Ala Gln Cys Val Leu Ala Glu Ile Pro

500 505 510

Gln Gln Val Val Gly Tyr Phe Asn Thr Tyr Lys Leu Leu Pro Pro Lys 515 520 525

Asn Pro Ala Thr Lys Gln Gln Lys Gln 530 535

<210> 254

<211> 537

<212> PRT

<213> Homo sapiens

<400> 254

Met Ala His Cys Val Thr Leu Val Gln Leu Ser Ile Ser Cys Asp His 1 5 10 15

Leu Ile Asp Lys Asp Ile Gly Ser Lys Ser Asp Pro Leu Cys Val Leu
20 25 30

Leu Gln Asp Val Gly Gly Ser Trp Ala Glu Leu Gly Arg Thr Glu
35 40 45

Arg Val Arg Asn Cys Ser Ser Pro Glu Phe Ser Lys Thr Leu Gln Leu 50 55 60

Glu Tyr Arg Phe Glu Thr Val Gln Lys Leu Arg Phe Gly Ile Tyr Asp
65 70 75 80

Ile Asp Asn Lys Thr Pro Glu Leu Arg Asp Asp Phe Leu Gly Gly
85 90 95

Ala Glu Cys Ser Leu Gly Gln Ile Val Ser Ser Gln Val Leu Thr Leu
100 105 110

Pro Leu Met Leu Lys Pro Gly Lys Pro Ala Gly Arg Gly Thr Ile Thr 115 120 125

Val Ser Ala Gln Glu Leu Lys Asp Asn Arg Val Val Thr Met Glu Val
130 135 140

Glu Ala Arg Asn Leu Asp Lys Lys Asp Phe Leu Gly Lys Ser Asp Pro 145 150 155 160

Phe Leu Glu Phe Phe Arg Gln Gly Asp Gly Lys Trp His Leu Val Tyr 165 170 175

| Arg | Ser | Glu | Val 180 | Ile | Lys | Asn | Asn | Leu 185 | Asn | Pro | Thr | Trp | Lys 190 | Arg | Phe |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser | Val | Pro 195 | Val | Gln | His | Phe | Cys 200 | Gly | Gly | Asn | Pro | Ser 205 | Thr | Pro | Ile |
| Gln | Val 210 | Gln | Cys | Ser | Asp | Tyr 215 | Asp | Ser | Asp | Gly | Ser 220 | His | Asp | Leu | Ile |
| Gly 225 | Thr | Phe | His | Thr | Ser 230 | Leu | Ala | Gln | Leu | Gln 235 | Ala | Val | Pro | Ala | Glu 240 |
| Phe | Glu | Cys | Ile | His 245 | Pro | Glu | Lys | Gln | Gln 250 | Lys | Lys | Lys | Ser | Tyr 255 | Lŷs |
| Asn | Ser | Gly | Thr 260 | Ile | Arg | Val | Lys | Ile 265 | Cys | Arg | Val | Glu | Thr 270 | Glu | Tyr |
| Ser | Phe | Leu 275 | Asp | Tyr | Val | Met | Gly 280 | Gly | Cys | Gln | Ile | Asn 285 | Phe | Thr | Val |
| Gly | Val 290 | Asp | Phe | Thr | Gly | Ser 295 | Asn | Gly | Asp | Pro | Ser 300 | Ser | Pro | Asp | Ser |
| Leu 305 | His | Tyr | Leu | Ser | Pro 310 | Thr | Gly | Val | Asn | Glu 315 | Tyr | Leu | Met | Ala | Leu 320 |
| Trp | Ser | Val | Gly | Ser 325 | Val | Val | Gln | Asp | Tyr 330 | Asp | Ser | Asp | Lys | Leu 335 | Phe |
| Pro | Ala | Phe | Gly 340 | Phe | Gly | Ala | Gln | Val 345 | Pro | Pro | Asp | Trp | Gln 350 | Val | Ser |
| His | Glu | Phe 355 | Ala | Leu | Asn | Phe | Asn 360 | Pro | Ser | Asn | Pro | Tyr 365 | Cys | Ala | Gly |
| Ile | Gln 370 | Gly | Ile | Val | Asp | Ala 375 | Tyr | Arg | Gln | Ala | Leu 380 | Pro | Gln | Val | Arg |
| Leu 385 | Tyr | Gly | Pro | Thr | Asn 390 | Phe | Ala | Pro | Ile | Ile 395 | Asn | His | Val | Ala | Arg 400 |
| Phe | Ala | Ala | Gln | Ala 405 | Ala | His | Gln | Gly | Thr 410 | Ala | Ser | Gln | Tyr | Phe 415 | Met |
| Leu | Leu | Leu | Leu 420 | Thr | Asp | Gly | Ala | Val 425 | Thr | Asp | Val | Glu | Ala | Thr | Arg |

Glu Ala Val Val Arg Ala Ser Asn Leu Pro Met Ser Val Ile Ile Val Gly Val Gly Gly Ala Asp Phe Glu Ala Met Glu Gln Leu Asp Ala Asp Gly Gly Pro Leu His Thr Arg Ser Gly Gln Ala Ala Ala Arg Asp Ile Val Gln Phe Val Pro Tyr Arg Phe Gln Asn Ala Pro Arg Glu Ala Leu Ala Gln Thr Val Leu Ala Glu Val Pro Thr Gln Leu Val Ser Tyr Phe Arg Ala Gln Gly Trp Ala Pro Leu Lys Pro Leu Pro Pro Ser Ala Lys Asp Pro Ala Gln Ala Pro Gln Ala <210> 255 <211> 454 <212> PRT <213> Mus musculus <400> 255 Met Ala His Cys Val Thr Leu Val Gln Leu Ser Val Ser Cys Glu His Leu Ile Asp Lys Asp Ile Gly Ser Lys Ser Asp Pro Leu Cys Val Leu 20 ' Leu Gln Asp Val Gly Gly Ala Trp Ala Glu Leu Cys Arg Thr Glu Arg Val Arg Asn Cys Ser Ser Pro Glu Phe Ser Lys Thr Leu Gln Ile Glu Tyr His Phe Glu Thr Val Gln Lys Leu Arg Phe Gly Ile Tyr Asp Ile Asp Asn Lys Thr Pro Glu Leu Gly Asp Asp Phe Leu Gly Gly Ala

Glu Cys Ser Leu Gly Gln Ile Val Ser Ser Gln Thr Leu Thr Leu Pro

| Leu Met I | Leu Lys 115 | Pro Gly | Lys | Pro 120 | Ala | Gly | Arg | Gly | Thr 125 | Ile | Thr | Val |
|------------------|----------------|----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser Ala (| Gln Glu | Leu Lys | Asp 135 | Ser | Arg | Val | Val | Thr 140 | Met | Glu | Val | Glu |
| Ala Arg A | Asn Leu | Asp Lys 150 | Lys | Asp | Phe | Leu | Gly 155 | Lys | Ser | Asp | Pro | Phe 160 |
| Leu Glu F | Phe Phe | Arg Gln 165 | Gly | Asp | Gly | Lys 170 | Trp | Gln | Leu | Ala | Tyr 175 | Arg |
| Thr Glu \ | Val Val 180 | Lys Asn | Asn | Leu | Asn 185 | Pro | Thr | Trp | Lys | Arg 190 | Phe | Ser |
| Val Ser I | Leu Gln 195 | His Phe | Cys | Gly 200 | Gly | Asp | Leu | Ser | Thr 205 | Pro | Ile | Gln |
| Val Arg (210 | Cys Ser | Asp Tyr | Asp 215 | Ser | Asp | Gly | Ser | His 220 | Asp | Leu | Ile | Gly |
| Thr Phe B | His Thr | Thr Leu 230 | Ala | Gln | Leu | Gln | Ala 235 | Val | Pro | Ala | Glu | Phe 240 |
| Glu Cys \ | Val His | Pro Glu 245 | Lys | Gln | Gln | Arg 250 | Lys | Lys | Asn | Tyr | Arg 255 | Asn |
| Ser Gly 1 | Thr Val 260 | Arg Val | Lys | Thr | Cys 265 | Arg | Val | Glu | Thr | Glu 270 | Tyr | Ser |
| Phe Leu A | Asp Tyr 275 | Val Met | Gly | Gly 280 | Cys | Gln | Ile | Asn | Phe 285 | Thr | Val | Gly |
| Val Asp I 290 | Phe Thr | Gly Ser | Asn 295 | Gly | Asp | Pro | Ser | Ser 300 | Pro | Asp | Ser | Leu |
| His Tyr I 305 | Leu Ser | Pro Thr | Gly | Val | Asn | Glu | Tyr 315 | Leu | Thr | Ala | Leu | Trp 320 |
| Ser Val (| Gly Ser | Val Val 325 | Gln | Asp | Tyr | Asp 330 | Ser | Asp | Lys | Leu | Phe 335 | Pro |
| Ala Phe (| Gly Phe 340 | Gly Ala | Gln | Val | Pro 345 | Pro | Asp | Trp | Gln | Val 350 | Ser | His |
| Glu Phe A | Ala Leu 355 | Asn Phe | Asn | Pro 360 | Ser | Asn | Pro | Tyr | Cys 365 | Ala | Gly | Ile |

Gln Gly Ile Val Asp Ala Tyr Arg Gln Ala Leu Pro Gln Val Arg Leu 370 375 380

Tyr Gly Pro Thr Asn Phe Ala Pro Ile Ile Asn His Val Ala Arg Phe 385 390 395 400

Ala Ala Gln Ala Gln Gln Arg Ser Ala Ser Gln Tyr Phe Val Leu 405 410 415

Leu Leu Thr Asp Gly Ala Val Thr Asp Val Glu Ala Thr Cys Lys
420 425 430

Ala Val Val Asp Ala Ser Lys Leu Pro Met Ser Val Ile Ile Val Gly
435 440 445

Val Gly Gly His Ser 450

<210> 256

<211> 94

<212> PRT

<213> Homo sapiens

<400> 256

Leu Ala Gly Arg Arg Leu Asp Lys Lys Asp Leu Phe Gly Lys Ser Asp
1 5 10 15

Pro Phe Leu Glu Phe Tyr Lys Pro Gly Asp Asp Gly Lys Trp Met Leu 20 · 25 30

Val His Arg Thr Glu Val Ile Lys Tyr Thr Leu Asp Pro Val Trp Lys
35 40 45

Pro Phe Thr Val Pro Leu Val Ser Leu Cys Asp Gly Asp Met Glu Lys 50 55 60

Pro Ile Gln Val Met Cys Tyr Asp Tyr Asp Asn Asp Gly Gly His Asp 65 70 75 80

Phe Ile Gly Glu Phe Gln Thr Ser Val Ser Gln Met Cys Glu $85 \hspace{1cm} 90$

<210> 257

<211> 88

<212> PRT

<220> <223> Description of Artificial Sequence: Protein kinase C conserved region 2 domain sequence <400> 257 Ile Ser Ala Arg Asn Leu Pro Pro Lys Asp Lys Gly Gly Lys Ser Asp 5 10 15 Pro Tyr Val Lys Val Ser Leu Asp Gly Asp Pro Arg Glu Lys Lys 25 20 30 Thr Lys Val Val Lys Asn Thr Leu Asn Pro Val Trp Asn Glu Thr Phe 40 45 Glu Phe Glu Val Pro Pro Pro Glu Leu Ser Glu Leu Glu Ile Glu Val 50 55 60 Tyr Asp Lys Asp Arg Phe Ser Arg Asp Asp Phe Ile Gly Arg Val Thr 70 75 65 Ile Pro Leu Ser Asp Leu Leu Leu 85 <210> 258 <211> 100 <212> PRT <213> Homo sapiens <400> 258 Val Ser Gly Gln Asn Leu Leu Asp Arg Asp Val Thr Ser Lys Ser Asp 5 1 10 15 Pro Phe Cys Val Leu Phe Thr Glu Asn Asn Gly Arg Trp Ile Glu Tyr 20 25 30 Asp Arg Thr Glu Thr Ala Ile Asn Asn Leu Asn Pro Ala Phe Ser Lys 35 40 45 Lys Phe Val Leu Asp Tyr His Phe Glu Glu Val Gln Lys Leu Lys Phe 50 55 60 Ala Leu Phe Asp Gln Asp Lys Ser Ser Met Arg Leu Asp Glu His Asp 70 65 75 80

<213> Artificial Sequence

Phe Leu Gly Gln Phe Ser Cys Ser Leu Gly Thr Ile Val Ser Ser Lys

95

Lys Ile Thr Arg 100

<210> 259

<211> 94

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Protein kinase C conserved region 2 domain sequence

<400> 259

Ile Ser Ala Arg Asn Leu Pro Pro Lys Asp Lys Gly Gly Lys Ser Asp 1 5 10 15

Pro Tyr Val Lys Val Ser Leu Asp Gly Asp Pro Arg Glu Lys Lys Lys 20 25 30

Thr Lys Val Val Lys Asn Thr Leu Asn Pro Val Trp Asn Glu Thr Phe
35 40 45

Glu Phe Glu Val Pro Pro Pro Glu Leu Ser Glu Leu Glu Ile Glu Val
50 55 60

Tyr Asp Lys Asp Arg Phe Ser Arg Asp Asp Phe Ile Gly Arg Val Thr 65 70 75 80

Ile Pro Leu Ser Asp Leu Leu Gly Gly Arg His Glu Lys
85 90

<210> 260

<211> 85

<212> PRT

<213> Homo sapiens

<400> 260

Val Ser Gly Gln Asn Leu Leu Asp Arg Asp Val Thr Ser Lys Ser Asp
1 5 10 15

Pro Phe Cys Val Leu Phe Thr Glu Asn Asn Gly Arg Trp Ile Glu Tyr
20 25 30

Asp Arg Thr Glu Thr Ala Ile Asn Asn Leu Asn Pro Ala Phe Ser Lys

35 40 45

Lys Phe Val Leu Asp Tyr His Phe Glu Glu Val Gln Lys Leu Lys Phe 50 55 60

Ala Leu Phe Asp Gln Asp Lys Ser Ser Met Arg Leu Asp Glu His Asp 65 70 75 80

Phe Leu Gly Gln Phe

<210> 261

<211> 82

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: C2 domain sequence

<400> 261

Ile Ser Ala Arg Asn Leu Pro Lys Met Asp Met Asn Gly Leu Ser Asp

1 5 10 15

Pro Tyr Val Lys Val Asp Leu Asp Gly Asp Pro Lys Asp Thr Lys Lys
20 25 30

Phe Lys Thr Lys Thr Val Lys Lys Thr Leu Asn Pro Val Trp Asn Glu 35 40 45

Thr Phe Val Phe Glu Lys Val Pro Leu Pro Asp Leu Ala Ser Leu Arg
50 55 60

Phe Ala Val Tyr Asp Glu Asp Arg Phe Ser Arg Asp Asp Phe Ile Gly 65 70 75 80

Gln Val

<210> 262

<211> 85

<212> PRT

<213> Homo sapiens

<400> 262

Leu Ala Gly Arg Arg Leu Asp Lys Lys Asp Leu Phe Gly Lys Ser Asp

1 5 10 15

Pro Phe Leu Glu Phe Tyr Lys Pro Gly Asp Asp Gly Lys Trp Met Leu 20 25 30

Val His Arg Thr Glu Val Ile Lys Tyr Thr Leu Asp Pro Val Trp Lys
35 40 45

Pro Phe Thr Val Pro Leu Val Ser Leu Cys Asp Gly Asp Met Glu Lys 50 55 60

Pro Ile Gln Val Met Cys Tyr Asp Tyr Asp Asn Asp Gly Gly His Asp 65 70 75 80

Phe Ile Gly Glu Phe

<210> 263

<211> 82

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: C2 domain sequence

<400> 263

Ile Ser Ala Arg Asn Leu Pro Lys Met Asp Met Asn Gly Leu Ser Asp 1 5 10 15

Pro Tyr Val Lys Val Asp Leu Asp Gly Asp Pro Lys Asp Thr Lys Lys
20 25 30

Phe Lys Thr Lys Thr Val Lys Lys Thr Leu Asn Pro Val Trp Asn Glu 35 40 45

Thr Phe Val Phe Glu Lys Val Pro Leu Pro Asp Leu Ala Ser Leu Arg 50 55 60

Phe Ala Val Tyr Asp Glu Asp Arg Phe Ser Arg Asp Asp Phe Ile Gly 65 70 75 80

Gln Val

<210> 264

<211> 174 <212> PRT <400> 264 65

<213> Homo sapiens

Met Gly Thr Asn Glu Tyr Leu Ser Ala Ile Trp Ala Val Gly Gln Ile 10

Ile Gln Asp Tyr Asp Ser Asp Lys Met Phe Pro Ala Leu Gly Phe Gly 20 25

Ala Gln Leu Pro Pro Asp Trp Lys Val Ser His Glu Phe Ala Ile Asn 40

Phe Asn Pro Thr Asn Pro Phe Cys Ser Gly Val Asp Gly Ile Ala Gln 55

Ala Tyr Ser Ala Cys Leu Pro His Ile Arg Phe Tyr Gly Pro Thr Asn 70 75

Phe Ser Pro Ile Val Asn His Val Ala Arg Phe Ala Ala Gln Ala Thr 85 90 95

Gln Gln Arg Thr Ala Thr Gln Tyr Phe Ile Leu Leu Ile Ile Thr Asp 100 105 110

Gly Val Ile Ser Asp Met Glu Glu Thr Arg His Ala Val Val Gln Ala 115 120 125

Ser Lys Leu Pro Met Ser Ile Ile Ile Val Gly Val Gly Asn Ala Asp 130 135 140

Phe Ala Ala Met Glu Phe Leu Asp Gly Asp Ser Arg Met Leu Arg Ser 145 150 155 160

His Thr Gly Glu Glu Ala Ala Arg Asp Ile Val Gln Phe Val 165 170

<210> 265

<211> 166

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: von Willebrand factor (vWF) type A domain sequence

<400> 265 Met Gly Gly Asn Arg Phe Glu Leu Ala Lys Glu Phe Val Leu Lys Leu 10 5 15 Val Glu Gln Leu Asp Ile Gly Pro Asp Gly Asp Arg Val Gly Leu Val 25 20 Thr Phe Ser Ser Asp Ala Arg Val Leu Phe Pro Leu Asn Asp Ser Gln 35 40 45 Ser Lys Asp Ala Leu Leu Glu Ala Leu Ala Ser Leu Ser Tyr Ser Leu 55 Gly Gly Thr Asn Leu Gly Ala Ala Leu Glu Tyr Ala Leu Glu Asn 70 75 65 Leu Phe Ser Glu Ser Ala Gly Ser Arg Arg Gly Ala Pro Lys Val Leu 90

Ile Leu Ile Thr Asp Gly Glu Ser Asn Asp Gly Gly Glu Asp Ile Leu
100 105 110

Lys Ala Ala Lys Glu Leu Lys Arg Ser Gly Val Lys Val Phe Val Val 115 120 125

Gly Val Gly Asn Asp Val Asp Glu Glu Glu Leu Lys Lys Leu Ala Ser 130 135 140

Ala Pro Gly Gly Val Phe Val Val Glu Asp Leu Pro Ser Leu Leu Asp 145 150 155 160

Leu Leu Ile Asp Leu Leu 165

<210> 266

<211> 416

<212> PRT

<213> Homo sapiens

<400> 266

Met Leu Ala Leu Leu Val Leu Val Thr Val Ala Leu Ala Ser Ala His

1 5 10 15

His Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg Val Asn Val 20 25 30

Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala Ser Thr Thr

| 35 | 40 | 45 |
|----|----|----|
| 33 | 40 | 40 |

| Gln | Ile 50 | Asp | Phe | Trp | Lys | Pro 55 | Asp | Ser | Val | Thr | Gln 60 | Ile | Lys | Pro | His |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-------------|------------|------------|
| Ser 65 | Thr | Val | Asp | Phe | Arg 70 | Val | Lys | Ala | Glu | Asp 75 | Thr | Val | Thr | Val | Glu 80 |
| Asn | Val | Leu | Lys | Gln 85 | Asn | Glu | Leu | Gln | Tyr 90 | Lys | Val | Leu | Ile | Ser 95 | Asr |
| Leu | Arg | Asn | Val 100 | Val | Glu | Ala | Gln | Phe 105 | Asp | Ser | Arg | Val | Arg 110 | Ala | Thi |
| Gly | His | Ser 115 | Tyr | Glu | Lys | Tyr | Asn 120 | Lys | Trp | Glu | Thr | Ile 125 | Glu | Ala | Trp |
| Thr | Gln 130 | Gln | Val | Ala | Thr | Glu 135 | Asn | Pro | Ala | Leu | Ile 140 | Ser | Arg | Ser | Val |
| Ile 145 | Gly | Thr | Thr | Phe | Glu 150 | Gly | Arg | Ala | Ile | Tyr 155 | Leu | Leu | Lys | Val | Gly 160 |
| Lys | Ala | Gly | Gln | Asn 165 | Lys | Pro | Ala | Ile | Phe 170 | Met | Asp | Cys | Gly | Phe 175 | His |
| Ala | Arg | Glu | Trp 180 | Ile | Ser | Pro | Ala | Phe 185 | Cys | Gln | Trp | | Val .190 | Arg | Gli |
| Ala | Val | Arg 195 | Thr | Tyr | Gly | Arg | Glu 200 | Ile | Gln | Val | Thr | Glu 205 | Leu | Leu | Asp |
| Lys | Leu 210 | Asp | Phe | Tyr | Val | Leu 215 | Pro | Val | Leu | Asn | Ile 220 | Asp | Gly | Tyr | Ile |
| Tyr 225 | Thr | Trp | Thr | Lys | Ser 230 | Arg | Phe | Trp | Arg | Lys 235 | Thr | Arg | Ser | Thr | His |
| Thr | Gly | Ser | Ser | Ile 245 | Gly | Thr | Asp | Pro | Asn 250 | Arg | Asn | Phe | Asp | Ala 255 | Gl |
| Trp | Cys | Glu | Ile 260 | Gly | Ala | Ser | Arg | Asn 265 | Pro | Cys | Asp | Glu | Thr 270 | Tyr | Cys |
| Gly | Pro | Ala 275 | Ala | Glu | Ser | Glu | Lys 280 | Glu | Thr | Lys | Ala | Leu 285 | Ala | Asp | Phe |

Ile Arg Asn Lys Leu Ser Ser Ile Lys Ala Tyr Leu Thr Ile His Ser

290 295 300

Tyr Ser Gln Met Met Ile Tyr Pro Tyr Ser Tyr Ala Tyr Lys Leu Gly 305 310 315

Glu Asn Asn Ala Glu Leu Asn Ala Leu Ala Lys Ala Thr Val Lys Glu 325 330 335

Leu Ala Ser Leu His Gly Thr Lys Tyr Thr Tyr Gly Pro Gly Ala Thr 340 345 350

Thr Ile Tyr Pro Ala Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp Gln 355 360 365

Gly Ile Arg Tyr Ser Phe Thr Phe Glu Leu Arg Asp Thr Gly Arg Tyr 370 375 380

Gly Phe Leu Leu Pro Glu Ser Gln Ile Arg Ala Thr Cys Glu Glu Thr 385 390 395 400

Phe Leu Ala Ile Lys Tyr Val Ala Ser Tyr Val Leu Glu His Leu Tyr
405 410 415

<210> 267

<211> 417

<212> PRT

<213> Homo sapiens

<400> 267

Met Leu Ala Leu Leu Val Leu Val Thr Val Ala Leu Ala Ser Ala His

1 5 10 15

His Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg Val Asn Val $20 \\ 25 \\ 30$

Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala Ser Thr Thr 35 40 45

Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile Lys Pro His
50 55 60

Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val Thr Val Glu 65 70 75 80

| Asn | Val | Leu | Lys | Gln 85 | Asn | Glu | Leu | Gln | Tyr 90 | Lys | Val | Leu | Ile | Ser 95 | Asn |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|--------------------------|------------|------------|
| Leu | Arg | Asn | Val 100 | Val | Glu | Ala | Gln | Phe 105 | Asp | Ser | Arg | Val | Arg 110 | Ala | Thr |
| Gly | His | Ser 115 | Tyr | Glu | Lys | Tyr | Asn 120 | Lys | Trp | Glu | Thr | Ile 125 | Glu | Ala | Trp |
| Thr | Gln 130 | Gln | Val | Ala | Thr | Glu 135 | Asn | Pro | Ala | Leu | Ile 140 | Ser | Arg | Ser | Val |
| Ile 145 | Gly | Thr | Thr | Phe | Glu 150 | Gly | Arg | Ala | Ile | Tyr 155 | Leu | Leu | Lys | Val | Gly 160 |
| Lys | Ala | Gly | Gln | Asn 165 | Lys | Pro | Ala | Ile | Phe 170 | Met | Asp | Cys | Gly | Phe 175 | His |
| Ala | Arg | Glu | Trp 180 | Ile | Ser | Pro | Ala | Phe 185 | Cys | Gln | Trp | Phe | Val 1 ['] 90 | Arg | Glu |
| Ala | Val | Arg 195 | Thr | Tyr | Gly | Arg | Glu 200 | Ile | Gln | Val | Thr | Glu 205 | Leu | Leu | Asp |
| Lys | Leu 210 | Asp | Phe | Tyr | Val | Leu 215 | Pro | Val | Leu | Asn | Ile 220 | Asp | Gly | Tyr | Ile |
| Tyr 225 | Thr | Trp | Thr | Lys | Ser 230 | Arg | Phe | Trp | Arg | Lys 235 | Thr | Arg | Ser | Thr | His 240 |
| Thr | Gly | Ser | Ser | Cys 245 | Ile | Gly | Thr | Asp | Pro 250 | Asn | Arg | Asn | Phe | Asp 255 | Ala |
| Gly | Trp | Cys | Glu 260 | Ile | Gly | Ala | Ser | Arg 265 | Asn | Pro | Cys | Asp | Glu 270 | Thr | Tyr |
| Cys | Gly | Pro 275 | Ala | Ala | Glu | Ser | Glu 280 | Lys | Glu | Thr | Lys | Ala 285 | Leu | Ala | Asp |
| Phe | Ile 290 | Arg | Asn | Lys | Leu | Ser 295 | Ser | Ile | Lys | Ala | Tyr 300 | Leu | Thr | Ile | His |
| | | | | | | | | | | | | | | | |
| Ser 305 | Tyr | Ser | Gln | Met | Met 310 | Ile | Tyr | Pro | Tyr | Ser 315 | Tyr | Ala | Tyr | Lys | Leu 320 |

Glu Leu Ala Ser Leu His Gly Thr Lys Tyr Thr Tyr Gly Pro Gly Ala 340 345 350

Thr Thr Ile Tyr Pro Ala Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp 355 360 365

Gln Gly Ile Arg Tyr Ser Phe Thr Phe Glu Leu Arg Asp Thr Gly Arg 370 375 380

Tyr Gly Phe Leu Leu Pro Glu Ser Gln Ile Arg Ala Thr Cys Glu Glu 385 390 395 400

Thr Phe Leu Ala Ile Lys Tyr Val Ala Ser Tyr Val Leu Glu His Leu 405 410 415

Tyr

<210> 268

<211> 417

<212> PRT

<213> Homo sapiens

<400> 268

Met Leu Ala Leu Leu Val Leu Val Thr Val Ala Leu Ala Ser Ala His

1 5 10 15

His Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg Val Asn Val
20 25 30

Glu Asp Glu Asn His Ile Asn Ile Ile Arg Glu Leu Ala Ser Thr Thr 35 40 45

Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile Lys Pro His 50 55 60

Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Thr Val Thr Val Glu 65 70 75 80

Asn Val Leu Lys Gln Asn Glu Leu Gln Tyr Lys Val Leu Ile Ser Asn 85 90 95

Leu Arg Asn Val Val Glu Ala Gln Phe Asp Ser Arg Val Arg Ala Thr
100 105 110

Gly His Ser Tyr Glu Lys Tyr Asn Lys Trp Glu Thr Ile Glu Ala Trp
115 120 125

| Thr | Gln 130 | Gln | Val | Ala | Thr | Glu 135 | Asn | Pro | Ala | Leu | Ile 140 | Ser | Arg | Ser | Val |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile 145 | Gly | Thr | Thr | Phe | Glu 150 | Gly | Arg | Ala | Ile | Tyr 155 | Leu | Leu | Lys | Val | Gly 160 |
| Lys | Ala | Gly | Gln | Asn 165 | Lys | Pro | Ala | Ile | Phe 170 | Met | Asp | Cys | Gly | Phe 175 | His |
| Ala | Arg | Glu | Trp 180 | Ile | Ser | Pro | Ala | Phe 185 | Cys | Gln | Trp | Phe | Val 190 | Arg | Glu |
| Ala | Val | Arg 195 | Thr | Tyr | Gly | Arg | Glu 200 | Ile | Gln | Val | Thr | Glu 205 | Leu | Leu | Asn |
| Lys | Leu 210 | Asp | Phe | Tyr | Val | Leu 215 | Pro | Val | Leu | Asn | Ile 220 | Asp | Gly | Tyr | Ile |
| Tyr 225 | Thr | Trp | Thr | Lys | Ser 230 | Arg | Phe | Trp | Arg | Lys 235 | Thr | Arg | Ser | Thr | His 240 |
| Thr | Gly | Ser | Ser | Cys 245 | Ile | Gly | Thr | Asp | Pro 250 | Asn | Arg | Asn | Phe | Asp 255 | Ala |
| Gly | Trp | Cys | Glu 260 | Ile | Gly | Ala | Ser | Arg 265 | Asn | Pro | Cys | Asp | Glu 270 | Thr | Tyr |
| Cys | Gly | Pro 275 | Ala | Ala | Glu | Ser | Glu 280 | Lys | Glu | Thr | Lys | Ala 285 | Leu | Ala | Asp |
| Phe | Ile 290 | Arg | Asn | Lys | Leu | Ser 295 | Ser | Ile | Lys | Ala | Tyr 300 | Leu | Thr | Ile | His |
| Ser 305 | Tyr | Ser | Gln | Met | Met 310 | Ile | Tyr | Pro | Tyr | Ser 315 | Tyr | Ala | Tyr | Lys | Leu 320 |
| Gly | Glu | Asn | Asn | Ala 325 | Glu | Leu | Asn | Ala | Leu 330 | Ala | Lys | Ala | Thr | Val 335 | Lys |
| Glu | Leu | Ala | Ser 340 | Leu | His | Gly | Thr | Lys 345 | Tyr | Thr | Tyr | Gly | Pro 350 | Gly | Ala |
| Thr | Thr | Ile 355 | Tyr | Pro | Ala | Ala | Gly 360 | Gly | Ser | Asp | Asp | Trp 365 | Ala | Tyr | Asp |
| Gln | Gly 370 | Ile | Arg | Tyr | Ser | Phe 375 | Thr | Phe | Glu | Leu | Arg 380 | Asp | Thr | Gly | Arg |

Tyr Gly Phe Leu Leu Pro Glu Ser Gln Ile Arg Ala Thr Cys Glu Glu 385 390 395 400

Thr Phe Leu Ala Ile Lys Tyr Val Ala Ser Tyr Val Leu Glu His Leu 405 410 415

Tyr

<210> 269

<211> 416

<212> PRT

<213> Sus scrofa

<400> 269

Met Leu Ala Phe Leu Ile Leu Val Thr Val Thr Leu Ala Ser Ala His
1 5 10 15

His Ser Gly Glu His Phe Glu Gly Glu Lys Val Phe Arg Val Asn Val 20 25 30

Glu Asp Glu Asn Asp Ile Ser Leu Leu His Glu Leu Ala Ser Thr Arg $35 \hspace{1cm} 40 \hspace{1cm} 45$

Gln Ile Asp Phe Trp Lys Pro Asp Ser Val Thr Gln Ile Lys Pro His
50 55 60

Ser Thr Val Asp Phe Arg Val Lys Ala Glu Asp Ile Leu Ala Val Glu 65 70 75 80

Asp Phe Leu Glu Gln Asn Glu Leu Gln Tyr Glu Val Leu Ile Asn Asn 85 90 95

Leu Arg Ser Val Leu Glu Ala Gln Phe Asp Ser Arg Val Arg Thr Thr
100 105 110

Gly His Ser Tyr Glu Lys Tyr Asn Asn Trp Glu Thr Ile Glu Ala Trp 115 120 125

Thr Lys Gln Val Thr Ser Glu Asn Pro Asp Leu Ile Ser Arg Thr Ala 130 135 140

Ile Gly Thr Thr Phe Leu Gly Asn Asn Ile Tyr Leu Leu Lys Val Gly
145 150 155 160

Lys Pro Gly Pro Asn Lys Pro Ala Ile Phe Met Asp Cys Gly Phe His

| Ala | Arg | Glu | Trp 180 | Ile | Ser | His | Ala | Phe 185 | Cys | Gln | Trp | Phe | Val 190 | Arg | Glu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Val | Leu 195 | Thr | Tyr | Gly | Tyr | Glu 200 | Ser | His | Met | Thr | Glu 205 | Phe | Leu | Asn |
| Lys | Leu 210 | Asp | Phe | Tyr | Val | Leu 215 | Pro | Val | Leu | Asn | Ile 220 | Asp | Gly | Tyr | Ile |
| Tyr 225 | Thr | Trp | Thr | Lys | Asn 230 | Arg | Met | Trp | Arg | Lys 235 | Thr | Arg | Ser | Thr | Asn 240 |
| Ala | Gly | Thr | Thr | Cys 245 | Ile | Gly | Thr | Asp | Pro 250 | Asn | Arg | Asn | Phe | Asp 255 | Ala |
| Gly | Trp | Cys | Thr 260 | Thr | Gly | Ala | Ser | Thr 265 | Asp | Pro | Cys | Asp | Glu 270 | Thr | Tyr |
| Cys | Gly | Ser 275 | Ala | Ala | Glu | Ser | Glu 280 | Lys | Glu | Thr | Lys | Ala 285 | Leu | Ala | Asp |
| Phe | Ile 290 | Arg | Asn | Asn | Leu | Ser 295 | Ser | Ile | Lys | Ala | Tyr 300 | Leu | Thr | Ile | His |
| Ser 305 | Tyr | Ser | Gln | Met | Ile 310 | Leu | Tyr | Pro | Tyr | Ser 315 | Tyr | Asp | Tyr | Lys | Leu 320 |
| Pro | Glu | Asn | Asn | Ala 325 | Glu | Leu | Asn | Asn | Leu 330 | Ala | Lys | Ala | Ala | Val 335 | Lys |
| Glu | Leu | Ala | Thr 340 | Leu | Tyr | Gly | Thr | Lys 345 | Tyr | Thr | Tyr | Gly | Pro 350 | Gly | Ala |
| Thr | Thr | Ile 355 | Tyr | Pro | Ala | Ala | Gly 360 | Gly | Ser | Asp | Asp | Trp 365 | Ala | Tyr | Asp |
| Gln | Gly 370 | Ile | Lys | Tyr | Ser | Phe 375 | Thr | Phe | Glu | Leu | Arg 380 | Asp | Lys | Gly | Arg |
| Tyr 385 | Gly | Phe | Ile | Leu | Pro 390 | Glu | Ser | Gln | Ile | Gln 395 | Ala | Thr | Cys | Glu | Glu 400 |
| Thr | Met | Leu | Ala | Ile 405 | Lys | Tyr | Val | Thr | Asn 410 | Tyr | Val | Leu | Gly | His 415 | Leu |

| <211 |)> 27 L> 41 | L6 | | | | | | | | | | | | | |
|------------|----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| | ?> PF 3> Ca | | fami | ilian | ris | | | | | | | | | | |
| <400 |)> 27 | 70 | | | | | | | | | | | | | |
| Met 1 | Ala | Phe | Leu | Ile 5 | Leu | Val | Thr | Leu | Ala 10 | Leu | Ala | Ser | Ala | His 15 | Tyr |
| Ser | Gly | Glu | His 20 | Phe | Glu | Gly | Glu | Lys 25 | Val | Phe | Arg | Val | Asn 30 | Val | Glu |
| Asp | Glu | Asn 35 | His | Ile | Asn | Leu | Leu 40 | His | Thr | Leu | Ala | Ser 45 | Thr | Thr | Gln |
| Ile | Asp 50 | Phe | Trp | Lys | Pro | Asp 55 | Ser | Val | Thr | Gln | Ile 60 | Lys | Pro | His | Ser |
| Thr 65 | Ala | Asp | Phe | Arg | Val 70 | Lys | Ala | Glu | Asp | Ile 75 | Leu | Thr | Val | Glu | Asp 80 |
| Phe | Leu | Lys | Gln | Asn 85 | Glu | Leu | His | Tyr | Glu 90 | Val | Leu | Ile | Asn | Asn 95 | Leu |
| Arg | Leu | Val | Leu 100 | Glu | Gly | Gln | Phe | Gly 105 | Arg | Gln | Val | Pr.o | Ala 110 | Thr | Gly |
| His | Ser | Tyr 115 | Glu | Lys | Tyr | Asn | Arg 120 | Trp | Glu | Thr | Ile | Glu 125 | Ala | Trp | Thr |
| Gln | Gln 130 | Val | Thr | Ser | Glu | Asn 135 | Pro | Asp | Leu | Ile | Ser 140 | Arg | Arg | Ser | Ile |
| Gly 145 | Thr | Thr | Phe | Glu | Gly 150 | Arg | Thr | Ile | Tyr | Leu 155 | Leu | Lys | Val | Gly | Lys 160 |
| Ala | Gly | Gln | Asn | Lys 165 | Pro | Ala | Ile | Phe | Met 170 | Asp | Cys | Gly | Phe | His 175 | Ala |
| Arg | Glu | Trp | Ile 180 | Ser | Pro | Ala | Phe | Trp 185 | Gln | Trp | Phe | Val | Arg 190 | Glu | Xaa |
| Ile | Arg | Thr | Tyr | Gly | Gln | Glu | Ile | His | Met | Thr | Glu | Leu | Leu | Asp | Lys |

| Leu | Asp 210 | Phe | Tyr | Val | Leu | Pro 215 | Val | Gly | Asn | Ile | Asp 220 | Gly | Tyr | Val | Tyr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Thr 225 | Trp | Thr | Lys | Asn | Arg 230 | Met | Trp | Arg | Lys | Thr 235 | Arg | Ser | Thr | Gln | Val 240 |
| Gly | Thr | Asn | Cys | Val 245 | Gly | Thr | Asp | Pro | Thr 250 | Arg | Asn | Phe | Asp | Ala 255 | Gly |
| Trp | Cys | Lys | Ile 260 | Gly | Ala | Ser | Arg | Asn 265 | Pro | Cys | Asp | | Thr 270 | Tyr | Cys |
| Gly | Pro | Ala 275 | Ala | Glu | Ser | Glu | Lys 280 | Glu | Thr | Lys | Ala | Leu 285 | Ala | Asn | Phe |
| Ile | Arg 290 | Ser | Asn | Leu | Ser | Ser 295 | Ile | Lys | Ala | Tyr | Leu 300 | Thr | Ile | His | Ser |
| Tyr 305 | Ser | Gln | Met | Met | Leu 310 | _ | Pro | Tyr | Ser | Tyr 315 | Asp | Tyr | Lys | Leu | Thr 320 |
| Glu | Asn | Asn | Ala | Glu 325 | Leu | Asn | Ala | Leu | Ala 330 | Lys | Ala | Thr | Val | Lys 335 | Glu |

Gly Ile Lys Tyr Ser Phe Thr Phe Glu Leu Arg Asp Lys Gly Arg Tyr 370 375 380

Leu Ala Thr Leu His Gly Thr Lys Tyr Thr Tyr Gly Pro Gly Ala Thr

Thr Ile Tyr Pro Ala Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp Gln

360

345

365

Gly Phe Ala Leu Pro Glu Ser Gln Ile Ser Pro Thr Cys Glu Glu Thr 385 390 395 400

Leu Leu Ala Ile Lys His Leu Ala Arg Tyr Val Leu Gln His Leu Tyr 405 410 415

<210> 271

<211> 82

<212> PRT

<213> Artificial Sequence

340

<220>

<223> Description of Artificial Sequence:Propep_M14 domain sequence

<400> 271

Gln Val Leu Arg Val Lys Val Ala Asp Glu Asp Gln Val Lys Leu Leu 1 5 10 15

Lys Asp Leu Glu Asn Thr Glu His Leu Glu Leu Asp Phe Trp Lys Pro 20 25 30

Asp Ser Ala Thr Pro Ile Lys Pro Gly Ser Thr Val Asp Phe Arg Val
35 40 45

Pro Ala Glu Asp Ile Gln Ala Val Lys Ser Phe Leu Glu Gln Ser Gly 50 55 60

Ile His Tyr Glu Val Leu Ile Glu Asp Val Gln Glu Leu Leu Glu Glu 65 70 75 80

Gln Phe

<210> 272

<211> 80

<212> PRT

<213> Homo sapiens

<400> 272

Lys Val Phe Arg Val Asn Val Glu Asp Glu Asn His Ile Asn Ile Ile 1 5 10 15

Arg Glu Leu Ala Thr Phe Ile Gln Ile Asp Phe Trp Lys Pro Asp Ser 20 25 30

Val Thr Gln Ile Lys Pro His Ser Thr Val Asp Phe Arg Val Lys Ala 35 40 45

Glu Asp Thr Val Thr Val Glu Asn Val Leu Lys Gln Asn Glu Leu Gln 50 55 60

Tyr Lys Val Leu Ile Ser Asn Leu Arg Asn Val Val Glu Ala Gln Phe 65 70 75 80

<213> Artificial Sequence <220> <223> Description of Artificial Sequence: Zn carbOpept domain sequence <400> 273 Tyr His Asn Leu Glu Glu Ile Tyr Ala Trp Leu Asp Leu Leu Val Ser Asn Phe Pro Asp Leu Val Ser Lys Val Ser Ile Gly Lys Ser Tyr Glu 20 Gly Arg Asp Leu Lys Val Leu Lys Ile Ser Asp Asn Pro Ala Thr Gly 35 40 Glu Asn Glu Pro Glu Val Phe Ala Val Ala Gly Trp Ile His Ala Arg 50 55 60 Glu Trp Val Thr Ser Ala Thr Leu Leu Trp Leu Leu Lys Glu Leu Val 65 70 75 80 Ala Asn Tyr Gly Ser Asp Lys Thr Ile Thr Lys Leu Leu Asp Gly Leu 85 90 Asp Leu Phe Tyr Ile Leu Pro Val Phe Asn Pro Asp Gly Tyr Ala Tyr 100 105 110 Ser Ile Thr Thr Asp Ser Tyr Arg Met Trp Arg Lys Thr 120 115 125 <210> 274 <211> 118 <212> PRT <213> Homo sapiens <400> 274 Tyr Asn Lys Trp Glu Thr Ile Glu Ala Trp Thr Gln Gln Val Ala Thr 10 15 Glu Asn Pro Ala Leu Ile Ser Arg Ser Val Ile Gly Thr Thr Phe Glu 20 25 30

<210> 273 <211> 125 <212> PRT Gly Arg Ala Ile Tyr Leu Leu Lys Val Gly Lys Ala Gly Gln Asn Lys
35 40 45

Pro Ala Ile Phe Met Glu Cys Gly Phe His Ala Arg Glu Trp Ile Ser 50 55 60

Pro Ala Phe Cys Gln Trp Phe Val Arg Glu Ala Val Arg Thr Tyr Gly 65 70 75 80

Arg Glu Ile Gln Val Thr Glu Leu Leu Asp Lys Leu Asp Phe Tyr Val 85 90 95

Leu Pro Val Leu Asn Ile Asp Gly Tyr Ile Tyr Thr Trp Thr Lys Ser 100 105 110

Arg Phe Trp Arg Lys Thr 115

<210> 275

<211> 101

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Zn_carbOpept
 domain sequence

<400> 275

Leu Leu Tyr Pro Tyr Gly Tyr Asp Tyr Asn Leu Asn Pro Asp Ala Asn 1 5 10 15

Asp Leu Asp Glu Leu Ser Asp Leu Lys Ile Ala Ala Asp Ala Leu Ser 20 25 30

Ala Arg His Gly Thr Tyr Tyr Thr Leu Gly Leu Pro Gly Ser Ser Thr
35 40 45

Ile Tyr Pro Ala Ser Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp Val
50 55 60

Gly Ile Ile Lys Tyr Ala Phe Thr Phe Glu Leu Arg Pro Asp Thr Gly
65 70 75 80

Ser Tyr Gly Asn Pro Cys Phe Leu Leu Pro Glu Glu Gln Ile Ile Pro 85 90 95

Thr Gly Ser Glu Glu

<210> 276 <211> 91 <212> PRT <213> Homo sapiens <400> 276 Trp Ile Tyr Pro Tyr Ser Tyr Ala Tyr Lys Leu Gly Glu Asn Asn Ala Glu Leu Asn Ala Leu Ala Lys Ala Thr Val Lys Glu Leu Ala Ser Leu His Gly Thr Lys Tyr Thr Tyr Gly Pro Gly Ala Thr Thr Ile Tyr Pro Ala Ala Gly Gly Ser Asp Asp Trp Ala Tyr Asp Gln Gly Ile Arg Tyr Ser Phe Thr Phe Glu Leu Arg Asp Thr Gly Arg Tyr Gly Phe Leu Leu Pro Glu Ser Gln Ile Arg Ala Thr Cys Glu Glu <210> 277 <211> 159 <212> PRT <213> Homo sapiens <400> 277 Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Arg Phe Ala Lys Lys His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asp Arg Leu 85 90 95

Ala Tyr Ile Ala His Pro Lys Leu Gly Lys Arg Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys
115 120 125

Ala Lys Asp Gln Thr Lys Ala Gln Ala Ala Ala Pro Ala Ser Val Pro 130 135 140

Ala Gln Ala Pro Lys Arg Thr Gln Ala Pro Thr Lys Ala Ser Glu 145 150 155

<210> 278

<211> 157

<212> PRT

<213> Homo sapiens

<400> 278

Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp

1 5 10 15

His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu 20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Arg Phe Ala Lys Lys
35 40 45

His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala 50 55 60

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Cys Lys Leu Asp Arg His
85 90 95

Ala Tyr Val Ala His Pro Lys Leu Gly Lys Arg Ala Leu Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys 115 120 125

Asp Gln Thr Lys Ala Gln Ala Ala Pro Ala Ser Val Pro Ala Gln

130 135 140

Ala Pro Lys Gly Thr Gln Ala Pro Thr Lys Ala Ser Glu 145 150 155

<210> 279

<211> 155

<212> PRT

<213> Homo sapiens

<400> 279

Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp

1 5 10 15

His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu 20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met His Phe Ala Lys Lys
35 40 45

His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala 50 55 60

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asp Arg Leu 85 90 95

Ala Tyr Ile Ala His Pro Lys Leu Gly Lys Arg Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys Asp Gln
115 120 125

Thr Lys Ala Gln Ala Ala Pro Pro Ser Val Pro Ala Gln Ala Pro 130 135 140

Lys Gly Ala Gln Ala Pro Thr Lys Ala Ser Glu 145 150 155

<210> 280

<211> 159

<212> PRT

<213> Homo sapiens

<400> 280

Met Ala Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp

1 5 10 15

His Arg Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu 20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Arg Phe Ala Lys Lys 35 40 45

His Asn Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Asn Ala Lys Ala 50 55 60

Met Ser Ala Arg Ala Glu Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asp Arg Leu 85 90 95

Ala Tyr Ile Ala His Pro Lys Leu Gly Lys Arg Ala Arg Ala Arg Ile 100 105 110

Ala Lys Gly Leu Arg Leu Cys Ala Pro Lys Ala Lys Ala Lys 115 120 125

Ala Lys Asp Gln Thr Lys Ala Gln Ala Ala Ala Pro Ala Ser Val Pro 130 135 140

Ala Gln Ala Pro Lys Arg Thr Gln Ala Pro Thr Lys Ala Ser Glu 145 150 155

<210> 281

<211> 189

<212> PRT

<213> Homo sapiens

<400> 281

Met Ala Lys Ser Lys Asn His Asn Thr His Asp Gln Phe Gln Lys Arg

1 5 10 15

His Arg Asn Gly Ile Lys Lys Pro Gln Ser Gln Arg Ser Val Ser Leu 20 25 30

Lys Gly Val Asp Pro Lys Phe Leu Arg Asn Met Pro Phe Ala Lys Lys 35 40 45

His Ser Lys Lys Gly Leu Lys Lys Met Gln Ala Asn Ser Ala Lys Ala

50 55 60

Met Ser Ala Arg Ala Lys Ala Ile Lys Ala Leu Val Lys Pro Lys Glu 65 70 75 80

Val Lys Pro Lys Ile Pro Lys Gly Val Ser Arg Lys Leu Asn Gln Leu 85 90 95

Ala Tyr Thr Gly Tyr Pro Lys Leu Gly Lys His Ala Cys Ala Arg Ile 100 105 110

Ala Lys Ala Leu Arg Leu Cys Arg Pro Lys Ala Lys Ala Lys Asp Gln 115 120 125

Thr Lys Ala Gln Ala Ala Ala Pro Ala Ser Val Pro Ala Gln Ala Pro 130 135 140

Lys Gly Ala Gln Ser Pro Tyr Lys Gly Phe Arg Val Glu Ile Ser Val 145 150 155 160

Cys Gln Arg Glu Asp Arg Arg Thr Gly Ala Thr Pro Pro Gly Cys His 165 170 175

Arg His Gly Ala Gly Val Leu Leu Cys Tyr Leu Tyr Lys 180 185

<210> 282

<211> 40

<212> PRT

<213> Homo sapiens

<400> 282

Lys Ser Lys Asn His Thr Thr His Asn Gln Ser Arg Lys Trp His Arg

1 5 10 15

Asn Gly Ile Lys Lys Pro Arg Ser Gln Arg Tyr Glu Ser Leu Lys Gly
20 25 30

Val Asp Pro Lys Phe Leu Arg Asn 35 40

<210> 283

<211> 40

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Ribosomal_L29e domain sequence

<400> 283

Lys Ser Lys Asn His Thr Asn His Asn Gln Asn Lys Lys Ala His Arg

1 5 10 15

Asn Gly Ile Lys Lys Pro Gln Lys Lys Arg Tyr Leu Ser Leu Lys Gly
20 25 30

Val Asp Ala Lys Phe Arg Arg Asn 35 40

<210> 284

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: tandem repeat unique to hRPL29

<400> 284

Lys Ala Lys Ala Lys Ala Lys Ala 1 5

<210> 285

<211> 790

<212> PRT

<213> Homo sapiens

<400> 285

Met Arg Ser Val Gln Ile Phe Leu Ser Gln Cys Arg Leu Leu Leu 1 5 10 15

Leu Val Pro Thr Met Leu Leu Lys Ser Leu Gly Glu Asp Val Ile Phe
20 25 30

His Pro Glu Gly Glu Phe Asp Ser Tyr Glu Val Thr Ile Pro Glu Lys
35 40 45

Leu Ser Phe Arg Gly Glu Val Gln Gly Val Val Ser Pro Val Ser Tyr
50 55 60

Leu Leu Gln Leu Lys Gly Lys Lys His Val Leu His Leu Trp Pro Lys

- Arg Leu Leu Pro Arg His Leu Arg Val Phe Ser Phe Thr Glu His
 85 90 95
- Gly Glu Leu Leu Glu Asp His Pro Tyr Ile Pro Lys Asp Cys Asn Tyr 100 105 110
- Met Gly Ser Val Lys Glu Ser Leu Asp Ser Lys Ala Thr Ile Ser Thr 115 120 125
- Cys Met Gly Gly Leu Arg Gly Val Phe Asn Ile Asp Ala Lys His Tyr 130 135 140
- Leu Leu Lys Lys Glu Gln Phe Gly Asn Gln Val Cys Gly Leu Ser Asp 165 170 175
- Asp Glu Ile Glu Trp Gln Met Ala Pro Tyr Glu Asn Lys Ala Arg Leu 180 185 190
- Arg Asp Phe Pro Gly Ser Tyr Lys His Pro Lys Tyr Leu Glu Leu Ile 195 200 205
- Leu Leu Phe Asp Gln Ser Arg Tyr Arg Phe Val Asn Asn Asn Leu Ser 210 215 220
- Gln Val Ile His Asp Ala Ile Leu Leu Thr Gly Ile Met Asp Thr Tyr 225 230 235 240
- Phe Gln Asp Val Arg Met Arg Ile His Leu Lys Ala Leu Glu Val Trp 245 250 255
- Thr Asp Phe Asn Lys Ile Arg Val Gly Tyr Pro Glu Leu Ala Glu Val 260 265 270
- Leu Gly Arg Phe Val Ile Tyr Lys Lys Ser Val Leu Asn Ala Arg Leu 275 280 285
- Ser Ser Asp Trp Ala His Leu Tyr Leu Gln Arg Lys Tyr Asn Asp Ala 290 295 300
- Leu Ala Trp Ser Phe Gly Lys Val Cys Ser Leu Glu Tyr Ala Gly Ser 305 310 315 320
- Val Ser Thr Leu Leu Asp Thr Asn Ile Leu Ala Pro Ala Thr Trp Ser

| - 1 · · · · · | ~ 1 | - | ~ 1 | - 1 | ~ 1 | ^ | - | ~ 1 | |
|---------------|-----|---|-----|---------|---------|-------|-------|-----|--|
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |
| | | | | | | | | | |

Ala His Glu Leu Gly His Ala Val Gly Met Ser His Asp Glu Gln Tyr Cys Gln Cys Arg Gly Arg Pro Asn Cys Ile Met Gly Ser Gly Arg Thr Gly Phe Ser Asn Cys Ser Tyr Ile Ser Phe Phe Lys His Ile Ser Ser Gly Ala Thr Cys Leu Asn Asn Ile Pro Gly Leu Gly Tyr Val Leu Lys Arg Cys Gly Asn Lys Ile Val Glu Asp Asn Glu Glu Cys Asp Cys Gly Ser Thr Glu Glu Cys Gln Lys Asp Arg Cys Cys Gln Ser Asn Cys Lys Leu Gln Pro Gly Ala Asn Cys Ser Ile Gly Leu Cys Cys His Asp Cys Arg Phe Arg Pro Ser Gly Tyr Val Cys Arg Gln Glu Gly Asn Glu Cys Asp Leu Ala Glu Tyr Cys Asp Gly Asn Ser Ser Ser Cys Pro Asn Asp Val Tyr Lys Gln Asp Gly Thr Pro Cys Lys Tyr Glu Gly Arg Cys Phe Arg Lys Gly Cys Arg Ser Arg Tyr Met Gln Cys Gln Ser Ile Phe Gly Pro Asp Ala Met Glu Ala Pro Ser Glu Cys Tyr Asp Ala Val Asn Leu Ile Gly Asp Gln Phe Gly Asn Cys Glu Ile Thr Gly Ile Arg Asn Phe Lys Lys Cys Glu Ser Ala Asn Ser Ile Cys Gly Arg Leu Gln Cys Ile Asn Val Glu Thr Ile Pro Asp Leu Pro Glu His Thr Thr Ile Ile Ser

Thr His Leu Gln Ala Glu Asn Leu Met Cys Trp Gly Thr Gly Tyr His

Leu Ser Met Lys Pro Met Gly Ile Pro Asp Leu Gly Met Ile Asn Asp 595 600 605

Gly Thr Ser Cys Gly Glu Gly Arg Val Cys Phe Lys Lys Asn Cys Val 610 615 620

Asn Ser Ser Val Leu Gln Phe Asp Cys Leu Pro Glu Lys Cys Asn Thr 625 630 635 640

Arg Gly Val Cys Asn Asn Arg Lys Asn Cys His Cys Met Tyr Gly Trp 645 650 655

Ala Pro Pro Phe Cys Glu Glu Val Gly Tyr Gly Gly Ser Ile Asp Ser 660 665 670

Gly Pro Pro Gly Leu Leu Arg Gly Ala Ile Pro Ser Ser Ile Trp Val 675 680 685

Val Ser Ile Ile Met Phe Arg Leu Ile Leu Leu Ile Leu Ser Val Val 690 695 700

Phe Val Phe Phe Arg Gln Val Ile Gly Asn His Leu Lys Pro Lys Gln 705 710 715 720

Glu Lys Met Pro Leu Ser Lys Ala Lys Thr Glu Gln Glu Glu Ser Lys
725 730 735

Thr Lys Thr Val Gln Glu Glu Ser Lys Thr Lys Thr Gly Gln Glu Glu 740 745 750

Ser Glu Ala Lys Thr Gly Gln Glu Glu Ser Lys Ala Lys Thr Gly Gln 755 760 765

Glu Glu Ser Lys Ala Asn Ile Glu Ser Lys Arg Pro Lys Ala Lys Ser 770 775 780

Val Lys Lys Gln Lys Lys 785 790

<210> 286

<211> 781

<212> PRT

<213> Homo sapiens

<400> 286

| Met 1 | Arg | Ser | Val | Gln 5 | Ile | Phe | Leu | Ser | Gln 10 | Cys | Arg | Leu | Leu | Leu 15 | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Val | Pro | Thr 20 | Met | Leu | Leu | Lys | Ser 25 | Leu | Gly | Glu | Asp | Val 30 | Ile | Phe |
| His | Pro | Glu 35 | Gly | Glu | Phe | Asp | Ser 40 | Tyr | Glu | Val | Thr | Ile 45 | Pro | Glu | Lys |
| Leu | Ser 50 | Phe | Arg | Gly | Glu | Val 55 | Gln | Gly | Val | Val | Ser 60 | Pro | Val | Ser | Tyr |
| Leu 65 | Leu | Gln | Leu | Lys | Gly 70 | Lys | Lys | His | Val | Leu 75 | His | Leu | Trp | Pro | Lys 80 |
| Arg | Leu | Leu | Leu | Pro 85 | Arg | His | Leu | Arg | Val 90 | Phe | Ser | Phe | Thr | Glu 95 | His |
| Gly | Glu | Leu | Leu 100 | Glu | Asp | His | Pro | Tyr 105 | Ile | Pro | Lys | Asp | Cys 110 | Asn | Tyr |
| Met | Gly | Ser 115 | Val | Lys | Glu | Ser | Leu 120 | Asp | Ser | Lys | Ala | Thr 125 | Ile | Ser | Thr |
| Cys | Met 130 | Gly | Gly | Leu | Arg | Gly 135 | Val | Phe | Asn | Ile | Asp 140 | Ala | Lys | His | Tyr |
| Gln 145 | Ile | Glu | Pro | Leu | Lys 150 | Ala | Ser | Pro | Ser | Phe 155 | Glu | His | Val | Val | Tyr 160 |
| Leu | Leu | Lys | Lys | Glu 165 | Gln | Phe | Gly | Asn | Gln 170 | Val | Cys | Gly | Leu | Ser 175 | Asp |
| Asp | Glu | Ile | Glu 180 | Trp | Gln | Met | Ala | Pro 185 | Tyr | Glu | Asn | Lys | Ala 190 | Arg | Leu |
| Arg | Asp | Phe 195 | Pro | Gly | Ser | Tyr | Lys 200 | His | Pro | Lys | Tyr | Leu 205 | Glu | Leu | Ile |
| Leu | Leu 210 | Phe | Asp | Gln | Ser | Arg 215 | Tyr | Arg | Phe | Val | Asn 220 | Asn | Asn | Leu | Ser |
| Gln 225 | Val | Ile | His | Asp | Ala 230 | Ile | Leu | Leu | Thr | Gly 235 | Ile | Met | Asp | Thr | Tyr 240 |
| Phe | Gln | Asp | Val | Arg | Met | Arg | Ile | His | Leu 250 | Lys | Ala | Leu | Glu | Val | Trp |

Thr Asp Phe Asn Lys Ile Arg Val Gly Tyr Pro Glu Leu Ala Glu Val Leu Gly Arg Phe Val Ile Tyr Lys Lys Ser Val Leu Asn Ala Arg Leu Ser Ser Asp Trp Ala His Leu Tyr Leu Gln Arg Lys Tyr Asn Asp Ala Leu Ala Trp Ser Phe Gly Lys Val Cys Ser Leu Glu Tyr Ala Gly Ser Val Ser Thr Leu Leu Asp Thr Asn Ile Leu Ala Pro Ala Thr Trp Pro Ala His Glu Leu Gly His Ala Val Gly Met Ser His Asp Glu Gln Tyr Cys Gln Cys Arg Gly Arg Leu Asn Cys Ile Met Gly Ser Gly Arg Thr Gly Phe Ser Asn Cys Ser Tyr Ile Ser Phe Phe Lys His Ile Ser Ser Gly Ala Thr Cys Leu Asn Asn Ile Pro Gly Leu Gly Tyr Val Leu Lys Arg Cys Gly Asn Lys Ile Val Glu Asp Asn Glu Glu Cys Asp Cys Gly Ser Thr Glu Glu Cys Gln Lys Asp Arg Cys Cys Gln Ser Asn Cys Lys Leu Gln Pro Gly Ala Asn Cys Ser Ile Gly Leu Cys Cys His Asp Cys Arg Phe Arg Pro Ser Gly Tyr Val Cys Arg Gln Glu Gly Asn Glu Cys Asp Leu Ala Glu Tyr Cys Asp Gly Asn Ser Ser Ser Cys Pro Asn Asp Val Tyr Lys Gln Asp Gly Thr Pro Cys Lys Tyr Glu Gly Arg Cys Phe Arg Lys Gly Cys Arg Ser Arg Tyr Met Gln Cys Gln Ser Ile Phe Gly

| Pro | Asp | Ala 515 | Met | Glu | Ala | Pro | Ser 520 | Glu | Cys | Tyr | Asp | Ala 525 | Val | Asn | Leu |
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| Ile | Gly 530 | Asp | Gln | Phe | Gly | Asn 535 | Cys | Glu | Ile | Thr | Gly 540 | Ile | Arg | Asn | Phe |
| Lys 545 | Lys | Cys | Glu | Ser | Ala 550 | Asn | Ser | Ile | Суѕ | Gly 555 | Arg | Leu | Gln | Cys | Ile 560 |
| Asn | Val | Glu | Thr | Ile 565 | Pro | Asp | Leu | Pro | Glu 570 | His | Thr | Thr | Ile | Ile 575 | Ser |
| Thr | His | Leu | Gln 580 | Ala | Glu | Asn | Leu | Met 585 | Cys | Trp | Gly | Thr | Gly 590 | Tyr | His |
| Leu | Ser | Met 595 | Lys | Pro | Met | Gly | Ile 600 | Pro | Asp | Leu | Gly | Met 605 | Ile | Asn | Asp |
| Gly | Thr 610 | Ser | Cys | Gly | Glu | Gly 615 | Arg | Val | Суѕ | Phe | Lys 620 | Lys | Asn | Cys | Val |
| Asn 625 | Ser | Ser | Val | Leu | Gln 630 | Phe | Asp | Cys | Leu | Pro 635 | Glu | Lys | Cys | Asn | Thr 640 |
| Arg | Gly | Val | Cys | Asn 645 | Asn | Arg | Lys | Asn | Cys 650 | His | Cys | Met | Tyr | Gly 655 | Trp |
| Ala | Pro | Pro | Phe 660 | Cys | Glu | Glu | Val | Gly 665 | Tyr | Gly | Gly | Ser | Ile 670 | Asp | Ser |
| Gly | Pro | Pro 675 | Gly | Leu | Leu | Arg | Gly 680 | Ala | Ile | Pro | Ser | Ser 685 | Ile | Trp | Val |
| Val | Ser 690 | Ile | Ile | Met | Phe | Arg 695 | Leu | Ile | Leu | Leu | Ile 700 | Leu | Ser | Val | Val |
| Phe 705 | Val | Phe | Phe | Arg | Gln 710 | Val | Ile | Gly | Asn | His 715 | Leu | Lys | Pro | Lys | Gln 720 |
| Glu | Lys | Met | Pro | Leu 725 | Ser | Lys | Ala | Lys | Thr 730 | Glu | Gln | Glu | Glu | Ser 735 | Lys |
| Thr | Lys | Thr | Val 740 | Gln | Glu | Glu | Ser | Lys 745 | Thr | Lys | Thr | Gly | Gln 750 | Glu | Glu |
| Ser | Glu | Ala 755 | Lys | Thr | Gly | Gln | Glu 760 | Glu | Ser | Lys | Ala | Asn 765 | Ile | Glu | Ser |

Lys Arg Pro Lys Ala Lys Ser Val Lys Lys Gln Lys Lys 770 780

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35 40 45

Ile Pro Leu Lys Val Thr Ser Arg Ala Arg Gly Ala Lys Asn Ser Glu 50 55 60

Trp Leu Ser Tyr Ser Leu Val Phe Gly Gly Arg Arg His Val Val His
65 70 75 80

Met Arg Val Lys Lys Leu Leu Val Ser Thr His Ile Pro Val Leu Thr 85 90 95

Tyr Thr Glu Glu His Thr Pro Leu Ser Asp Tyr Pro Phe Val Pro Ser 100 105 110

Asp Cys Tyr Tyr His Gly Tyr Val Glu Gly Ala Leu Glu Ser Leu Val 115 120 125

Ala Phe Ser Ala Cys Asn Gly Gly Leu Gln Gly Val Leu Gln Met Asn 130 135 140

Gly Phe Ser Tyr Glu Ile Glu Pro Ile Lys His Ser Ser Thr Phe Glu 145 150 155 160

His Leu Val Tyr Thr Leu Asn Asn Asn Lys Thr Gln Phe Pro Pro Met 165 170 175

Leu Cys Ser Leu Thr Glu Lys Arg Leu Leu Tyr Gln Pro Phe Gly Val 180 185 190

Glu Glu Ala Lys Lys Ser Ala Met Lys Gln Asn Tyr Gly Lys Leu Trp 195 200 205

| Pro | His 210 | Met | Trp | Phe | Leu | Glu 215 | Leu | Ala | Val | Val | Val 220 | Asp | Tyr | Gly | Phe |
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| Phe 225 | Thr | Asn | Ala | Gln | Gln 230 | Asn | Leu | Ser | Lys | Val 235 | Arg | Gly | Asp | Val | Val 240 |
| Leu | Val | Val | Asn | Met 245 | Val | Asp | Ser | Met | Tyr 250 | Lys | Pro | Leu | Asp | Thr 255 | Tyr |
| Val | Thr | Leu | Val 260 | Gly | Ile | Glu | Ile | Trp 265 | Asn | Arg | Gly | Asn | Val 270 | Leu | Pro |
| Met | Glu | Asn 275 | Ile | His | Gln | Val | Leu 280 | Glu | Asp | Phe | Ser | His 285 | Trp | Lys | Gln |
| Ile | Ser 290 | Leu | Ser | Gln | Val | His 295 | His | Asp | Ala | Ala | His 300 | Ile | Phe | Ile | Arg |
| Ser 305 | Ser | Leu | Ile | Ser | Val 310 | Leu | Gly | Ile | Ala | Tyr _. 315 | Ile | Ala | Gly | Ile | Cys 320 |
| Arg | Pro | Pro | Leu | Asp 325 | Cys | Gly | Val | Glu | Asn 330 | Phe | Gln | Gly | Asp | Ala 335 | Trp |
| Ser | Leu | Phe | Ala 340 | Asn | Thr | Val | Ala | His 345 | Glu | Leu | Gly | His | Thr 350 | Phe | Gly |
| Met | Lys | His 355 | Asp | Glu | Glu | Ser | Cys 360 | Ser | Cys | Gly | Lys | Ser 365 | Gly | Cys | Val |
| Met | Ser 370 | Thr | Phe | Arg | Val | Pro 375 | Ala | Glu | Arg | Phe | Thr 380 | Asn | Cys | Ser | Tyr |
| Ser 385 | Asp | Phe | Met | Lys | Thr 390 | Thr | Leu | Asn | Gln | Gly 395 | Thr | Cys | Leu | Tyr | Asn 400 |
| His | Pro | Arg | Pro | Gly 405 | Ala | Gly | Phe | Leu | Val 410 | Lys | Arg | Cys | Gly | Asn 415 | Gly |
| Met | Val | Glu | Ser 420 | Glu | Glu | Glu | Cys | Asp 425 | Cys | Gly | Ser | Val | Gln 430 | Glu | Cys |
| Glu | Gln | Asp 435 | Pro | Cys | Cys | Phe | Leu 440 | Asn | Cys | Thr | Leu | Arg 445 | Pro | Ala | Ala |
| Ala | | | | | | | | | | | | | | | |

| Gly 465 | Glu | Leu | Cys | Arg | Pro 470 | Lys | Ile | Asn | Glu | Cys 475 | Asp | Leu | Pro | Glu | Trp 480 |
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| Cys | Asn | Gly | Thr | Ser 485 | His | Gln | Cys | Pro | Glu 490 | Asp | Gly | Tyr | Val | Gln 495 | Asp |
| Gly | Val | Pro | Cys 500 | Gly | Ala | Gly | Ala | Tyr 505 | Cys | Tyr | Gln | Lys | Gln 510 | Cys | Asn |
| Asn | His | Asp 515 | Gln | Gln | Cys | Arg | Glu 520 | Ile | Phe | Gly | Lys | Gly 525 | Ala | Arg | Ser |
| Ala | Ser 530 | His | Asn | Cys | Tyr | Lys 535 | Glu | Ile | Asn | Leu | Gln 540 | Gly | Asn | Arg | Phe |
| Gly 545 | His | Cys | Gly | Thr | Asp 550 | Gly | Thr | Val | Phe | Leu 555 | Lys | Cys | Arg | Met | Ser 560 |
| Asp | Val | Phe | Cys | Gly 565 | Lys | Val | His | Cys | Glu 570 | Asn | Val | Glu | Asp | Ile 575 | His |
| His | Pro | Gln | Ala 580 | Pro | Tyr | Val | Leu | Gln 585 | Asn | Ile | Tyr | Ala | Asn 590 | Gly | Ile |
| Thr | Cys | Trp 595 | Ser | Thr | Gly | His | Cys 600 | Leu | Gly | Met | Gly | Val 605 | Pro | Asp | Val |
| Gly | Glu 610 | Val | Lys | Asp | Gly | Thr 615 | Thr | Cys | Gly | Val | Gly 620 | Lys | Ile | Cys | Leu |
| His 625 | Lys | Lys | Cys | Val | Ser 630 | Leu | Ser | Val | Leu | Ser 635 | Asn | Ala | Cys | Leu | Pro 640 |
| Glu | Thr | Cys | Asn | Arg 645 | Lys | Gly | Val | Cys | Asn 650 | Asn | Lys | His | His | Cys 655 | His |
| Cys | Asp | Tyr | Gly 660 | Trp | Ser | Pro | Pro | Phe 665 | Cys | Leu | His | Arg | Gly 670 | Tyr | Gly |
| Gly | Ser | Ile 675 | Asp | Ser | Gly | Pro | Thr 680 | Ser | Gln | Lys | Arg | Arg 685 | Val | Ile | Ile |
| Thr | Val 690 | Leu | Ser | Ile | Thr | Val 695 | Pro | Val | Leu | Ser | Ile 700 | Leu | Ile | Cys | Leu |
| Leu 705 | Ile | Ala | Gly | Leu | Tyr 710 | Arg | Ile | Tyr | Cys | Lys 715 | Ile | Pro | Ser | Gly | Pro 720 |

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Ser Ala Leu Glu Pro Lys Ser Ala Gly Asp Trp Trp Thr His Ala Trp

195 200 205

| Phe | Leu 210 | Glu | Leu | Val | Val | Val 215 | Val | Asn | His | Asp | Phe 220 | Phe | Ile | Tyr | Ser |
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| Gln 225 | Ser | Asn | Ile | Ser | Lys 230 | Val | Gln | Glu | Asp | Val 235 | Phe | Leu | Val | Val | Asn 240 |
| Ile | Val | Asp | Ser | Met 245 | Tyr | Lys | Gln | Leu | Gly 250 | Thr | Tyr | Ile | Ile | Leu 255 | Ile |
| Gly | Ile | Glu | Ile 260 | Trp | Asn | Gln | Gly | Asn 265 | Val | Phe | Pro | Met | Thr 270 | Ser | Ile |
| Glu | Gln | Val 275 | Leu | Asn | Asp | Phe | Ser 280 | Gln | Trp | Lys | Gln | Ile 285 | Ser | Leu | Ser |
| Gln | Leu 290 | Gln | His | Asp | Ala | Ala 295 | His | Met | Phe | Ile | Lys 300 | Asn | Ser | Leu | Ile |
| Ser 305 | Ile | Leu | Gly | Leu | Ala 310 | Tyr | Val | Ala | Gly | Ile 315 | Cys | Arg | Pro | Pro | Ile 320 |
| Asp | Cys | Gly | Val | Asp 325 | Asn | Phe | Gln | Gly | Asp 330 | Thr | Trp | Ser | Leu | Phe 335 | Ala |
| Asn | Thr | Val | Ala 340 | His | Glu | Leu | Gly | His 345 | Thr | Leu | Gly | Met | Gln 350 | His | Asp |
| Glu | Glu | Phe 355 | Cys | Phe | Cys | Gly | Glu 360 | Arg | Gly | Cys | Ile | Met 365 | Asn | Thr | Phe |
| Arg | Val 370 | Pro | Ala | Glu | Lys | Phe 375 | Thr | Asn | Cys | Ser | Tyr 380 | Ala | Asp | Phe | Met |
| Lys 385 | Thr | Thr | Leu | Asn | Gln 390 | Gly | Ser | Cys | Leu | His 395 | Asn | Pro | Pro | Arg | Leu 400 |
| Gly | Glu | Ile | Phe | Met 405 | Leu | Lys | Arg | Cys | Gly 410 | Asn | Gly | Val | Val | Glu 415 | Arg |
| Glu | Glu | Gln | Cys 420 | Asp | Cys | Gly | Ser | Val 425 | Gln | Gln | Cys | Glu | Gln 430 | Asp | Ala |
| Cys | Cys | Leu 435 | Leu | Asn | Cys | Thr | Leu 440 | Arg | Pro | Gly | Ala | Ala 445 | Cys | Ala | Phe |
| Gly | Leu | Cys | Cys | Lys | Asp | Cys | Lys | Phe | Met | Pro | Ser | Gly | Glu | Leu | Cys |

| 450 | 455 | 460 |
|-----|-----|-----|
| | | |

| Arg 465 | Gln | Glu | Val | Asn | Glu 470 | Cys | Asp | Leu | Pro | Glu 475 | Trp | Cys | Asn | Gly | Thr 480 |
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| Ser | His | Gln | Cys | Pro 485 | Glu | Asp | Arg | Tyr | Val 490 | Gln | Asp | Gly | Ile | Pro 495 | Cys |
| Ser | Asp | Ser | Ala 500 | Tyr | Cys | Tyr | Gln | Lys 505 | Arg | Суѕ | Asn | Asn | His 510 | Asp | Gln |
| His | Cys | Arg 515 | Glu | Ile | Phe | Gly | Lys 520 | Asp | Ala | Lys | Ser | Ala 525 | Ser | Gln | Asn |
| Cys | Tyr 530 | Lys | Glu | Ile | Asn | Ser 535 | Gln | Gly | Asn | Arg | Phe 540 | Gly | His | Cys | Gly |
| Ile 545 | Asn | Gly | Thr | Thr | Tyr 550 | Leu | Lys | Cys | His | Ile 555 | Ser | Asp | Val | Phe | Cys 560 |
| Gly | Arg | Val | Gln | Cys 565 | Glu | Asn | Val | Arg | Asp 570 | Ile | Pro | Leu | Leu | Gln 575 | Asp |
| His | Phe | Thr | Leu 580 | Gln | His | Thr | His | Ile 585 | Asn | Gly | Val | Thr | Cys 590 | Trp | Gly |
| Ile | Asp | Tyr 595 | His | Leu | Arg | Met | Asn 600 | Ile | Ser | Asp | Ile | Gly 605 | Glu | Val | Lys |
| Asp | Gly 610 | Thr | Val | Cys | Gly | Pro 615 | Gly | Lys | Ile | Cys | Ile 620 | His | Lys | Lys | Cys |
| Val 625 | Ser | Leu | Ser | Val | Leu 630 | Ser | His | Val | Cys | Leu 635 | Pro | Glu | Thr | Cys | Asn 640 |
| Met | Lys | Gly | Ile | Cys 645 | Asn | Asn | Lys | His | His 650 | Cys | His | Cys | Gly | Tyr 655 | Gly |
| Trp | Ser | Pro | Pro 660 | Tyr | Cys | Gln | His | Arg 665 | Gly | Tyr | Gly | Gly | Ser 670 | Ile | Asp |
| Ser | Gly | Pro 675 | Ala | Ser | Ala | Lys | Arg 680 | Gly | Val | Phe | Leu | Pro 685 | Leu | Ile | Val |
| Ile | Pro 690 | Ser | Leu | Ser | Val | Leu 695 | Thr | Phe | Leu | Phe | Thr 700 | Val | Gly | Leu | Leu |
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Ile Ser Arg Gly Arg Ser Ala Lys Ala Pro Gly Trp Leu Ser Tyr Ser
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Leu Arg Phe Gly Gly Gln Lys His Val Val His Met Arg Val Lys Lys 65 70 75 80

Leu Leu Val Ser Arg His Leu Pro Val Phe Thr Tyr Thr Asp Glu Arg
85 90 95

Ala Leu Leu Glu Asp Gln Leu Phe Ile Pro Asp Asp Cys Tyr Tyr His 100 105 110

Gly Tyr Val Glu Gly Ala Pro Glu Ser Leu Val Val Phe Ser Ala Cys 115 120 125

Phe Gly Gly Phe Arg Gly Val Leu Lys Ile Ser Gly Leu Thr Tyr Glu 130 135 140

Val Asn Ser Asn Glu Thr Gln Phe Pro Ala Met Arg Cys Gly Leu Thr 165 170 175

Glu Lys Glu Val Ala Arg Gln Gln Leu Glu Phe Glu Glu Ala Glu Asn 180 185 190

| Ser | Ala | Leu 195 | Glu | Pro | Lys | Ser | Ala 200 | Gly | Asp | Trp | Trp | Thr 205 | His | Ala | Trp |
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| Phe | Leu 210 | Glu | Leu | Val | Val | Val 215 | Val | Asn | His | Asp | Phe 220 | Phe | Ile | Tyr | Ser |
| Gln 225 | Ser | Asn | Ile | Ser | Lys 230 | Val | Gln | Glu | Asp | Val 235 | Phe | Leu | Val | Val | Asn 240 |
| Ile | Val | Asp | Ser | Met 245 | Tyr | Gln | Gln | Leu | Gly 250 | Thr | Tyr | Ile | Ile | Leu 255 | Ile |
| Gly | Ile | Glu | Ile 260 | Trp | Asn | Gln | Gly | Asn 265 | Val | Phe | Pro | Met | Thr 270 | Ser | Ile |
| Glu | Gln | Val 275 | Leu | Asn | Asp | Phe | Ser 280 | Gln | Trp | Lys | Gln | Ile 285 | Ser | Leu | Ser |
| Gln | Leu 290 | Gln | His | Asp | Ala | Ala 295 | His | Met | Phe | Ile | Lys 300 | Asn | Ser | Leu | Ile |
| Ser 305 | Ile | Leu | Gly | Leu | Ala 310 | Tyr | Val | Ala | Gly | Ile 315 | Cys | Arg | Pro | Pro | Ile 320 |
| Asp | Cys | Gly | Val | Asp 325 | Asn | Phe | Gln | Gly | Asp 330 | Thr | Trp | Ser | Leu | Phe 335 | Ala |
| Asn | Thr | Val | Ala 340 | His | Glu | Leu | Gly | His 345 | Thr | Leu | Gly | Met | Gln 350 | His | Asp |
| Glu | Glu | Phe 355 | Cys | Phe | Cys | Gly | Glu 360 | Arg | Gly | Cys | Ile | Met 365 | Asn | Thr | Phe |
| Arg | Val 370 | Pro | Ala | Glu | Lys | Phe 375 | Thr | Asn | Cys | Ser | Tyr 380 | Ala | Asp | Phe | Met |
| Lys 385 | Thr | Thr | Leu | Asn | Gln 390 | Gly | Ser | Cys | Leu | His 395 | Asn | Pro | Pro | Arg | Leu 400 |
| Gly | Glu | Ile | Phe | Met 405 | Leu | Lys | Arg | Cys | Gly 410 | Asn | Gly | Val | Val | Glu 415 | Arg |
| Glu | Glu | Gln | Cys 420 | Asp | Cys | Gly | Ser | Val 425 | Gln | Gln | Cys | Glu | Gln 430 | Asp | Ala |
| Cys | Cys | Leu 435 | Leu | Asn | Cys | Thr | Leu 440 | Arg | Pro | Gly | Ala | Ala 445 | Суз | Ala | Phe |

| Gly | Leu 450 | Cys | Cys | Lys | Asp | Cys 455 | Lys | Phe | Met | Pro | Ser 460 | Gly | Glu | Leu | Cys |
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| Arg 465 | Gln | Glu | Val | Asn | Glu 470 | Cys | Asp | Leu | Pro | Glu 475 | Trp | Cys | Asn | Gly | Thr 480 |
| Ser | His | Gln | Cys | Pro 485 | Glu | Asp | Arg | Tyr | Val 490 | Gln | Asp | Gly | Ile | Pro 495 | Cys |
| Ser | Asp | Ser | Ala 500 | Tyr | Cys | Tyr | Gln | Lys 505 | Arg | Cys | Asn | Asn | His 510 | Asp | Gln |
| His | Cys | Arg 515 | Glu | Ile | Phe | Gly | Lys 520 | Asp | Ala | Lys | Ser | Ala 525 | Ser | Gln | Asn |
| Cys | Tyr 530 | Lys | Glu | Ile | Asn | Ser 535 | Gln | Gly | Asn | Arg | Phe 540 | Gly | His | Cys | Gly |
| Ile 545 | Asn | Gly | Thr | Thr | Tyr 550 | Leu | Lys | Cys | His | Ile 555 | Ser | Asp | Val | Phe | Cys 560 |
| Gly | Arg | Val | Gln | Cys 565 | Glu | Asn | Val | Arg | Asp 570 | Ile | Pro | Leu | Leu | Gln 575 | Asp |
| His | Phe | Thr | Leu 580 | Gln | His | Thr | His | Ile 585 | Asn | Gly | Val | Thr | Cys 590 | Trp | Gly |
| Ile | Asp | Tyr 595 | His | Leu | Arg | Met | Asn 600 | Ile | Ser | Asp | Ile | Gly 605 | Glu | Val | Lys |
| Asp | Gly 610 | Thr | Val | Cys | Gly | Pro 615 | Gly | Lys | Ile | Cys | Ile 620 | His | Lys | Lys | Cys |
| Val 625 | Ser | Leu | Ser | Val | Leu 630 | Ser | His | Val | Cys | Leu 635 | Pro | Glu | Thr | Cys | Asn 640 |
| Met | Lys | Gly | Ile | Cys 645 | Asn | Asn | Lys | His | His 650 | Cys | His | Cys | Gly | Tyr 655 | Gly |
| Trp | Ser | Pro | Pro 660 | Tyr | Cys | Gln | His | Arg 665 | Gly | Tyr | Gly | Gly | Ser 670 | Ile | Asp |
| Ser | Gly | Pro 675 | Ala | Ser | Ala | Lys | Arg 680 | Gly | Val | Phe | Leu | Pro 685 | Leu | Ile | Val |
| Ile | Pro 690 | Ser | Leu | Ser | Val | Leu 695 | Thr | Phe | Leu | Phe | Thr 700 | Val | Gly | Leu | Leu |

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Thr Tyr Asp Asp Gly Thr Leu Val Thr Glu His Pro Leu Ile Gln

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Ser Ile Asp Phe Lys Gln Val Leu His Ser Trp Phe Arg Gln Pro Gln

Leu Ala Val Thr Ser Leu Gly Pro Gly Ala Glu Gly Leu His Pro Phe 245 250 Met Glu Leu Arg Val Leu Glu Asn Thr Lys Arg Ser Arg Asn Leu 260 265 270 Gly Leu Asp Cys Asp Glu His Ser Ser Glu Ser Arg Cys Cys Arg Tyr 275 280 285 Pro Leu Thr Val Asp Phe Glu Ala Phe Gly Trp Asp Trp Ile Ile Ala 295 Pro Lys Arg Tyr Lys Ala Asn Tyr Cys Ser Gly Gln Cys Glu Tyr Met 310 315 Phe Met Gln Lys Tyr Pro His Thr His Leu Val Gln Gln Ala Asn Pro 325 330 Arg Gly Ser Ala Gly Pro Cys Cys Thr Pro Thr Lys Met Ser Pro Ile 345 Asn Met Leu Tyr Phe Asn Asp Lys Gln Gln Ile Ile Tyr Gly Lys Ile 355 360 365 Pro Gly Met Val Val Asp Arg Cys Gly Cys Ser 370 375 <210> 295 <211> 407 <212> PRT <213> Homo sapiens <400> 295 Met Val Leu Ala Ala Pro Leu Leu Gly Phe Leu Leu Leu Ala Leu 10 Glu Leu Arg Pro Arg Gly Glu Ala Ala Glu Gly Pro Ala Ala Ala Ala 20 25 30 Ala Ala Ala Ala Ala Ala Ala Ala Gly Val Gly Glu Arg Ser Ser Arg Pro Ala Pro Ser Val Ala Pro Glu Pro Asp Gly Cys Pro Val

Ser Asn Trp Gly Ile Glu Ile Asn Ala Phe Asp Pro Ser Gly Thr Asp

235

240

230

| Cys | Val | Trp | Arg | Gln | His | Ser | Arg | Glu | Leu | Arg | Leu | Glu | Ser | Ile | Lys |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| 65 | | | | | 70 | | | | | 75 | | | | | 80 |

60

55

- Ser Gln Ile Leu Ser Lys Leu Arg Leu Lys Glu Ala Pro Asn Ile Ser
- Arg Glu Val Val Lys Gln Leu Leu Pro Lys Ala Pro Pro Leu Gln Gln
 100 105 110
- Ile Leu Asp Leu His Asp Phe Gln Gly Asp Ala Leu Gln Pro Glu Asp 115 120 125
- Phe Leu Glu Glu Asp Glu Tyr His Ala Thr Thr Glu Thr Val Ile Ser 130 135 140
- Met Ala Gln Glu Thr Asp Pro Ala Val Gln Thr Asp Gly Ser Pro Leu 145 150 155 160
- Cys Cys His Phe His Phe Ser Pro Lys Val Met Phe Thr Lys Val Leu 165 170 175
- Lys Ala Gln Leu Trp Val Tyr Leu Arg Pro Val Pro Arg Pro Ala Thr 180 185 190
- Val Tyr Leu Gln Ile Leu Arg Leu Lys Pro Leu Thr Gly Glu Gly Thr 195 200 205
- Ala Gly Gly Gly Gly Gly Arg Arg His Ile Arg Ile Arg Ser Leu 210 215 220
- Lys Ile Glu Leu His Ser Arg Ser Gly His Trp Gln Ser Ile Asp Phe 225 230 235 240
- Lys Gln Val Leu His Ser Trp Phe Arg Gln Pro Gln Ser Asn Trp Gly
 245 250 255
- Ile Glu Ile Asn Ala Phe Asp Pro Ser Gly Thr Asp Leu Ala Val Thr 260 265 270
- Ser Leu Gly Pro Gly Ala Glu Gly Leu His Pro Phe Met Glu Leu Arg 275 280 285
- Val Leu Glu Asn Thr Lys Arg Ser Arg Arg Asn Leu Gly Leu Asp Cys 290 295 300
- Asp Glu His Ser Ser Glu Ser Arg Cys Cys Arg Tyr Pro Leu Thr Val

| 305 | 310 | 315 | 320 |
|-----|-----|-----|-----|
| | | | |

Asp Phe Glu Ala Phe Gly Trp Asp Trp Ile Ile Ala Pro Lys Arg Tyr 325 330 335

Lys Ala Asn Tyr Cys Ser Gly Gln Cys Glu Tyr Met Phe Met Gln Lys 340 345 350

Tyr Pro His Thr His Leu Val Gln Gln Ala Asn Pro Arg Gly Ser Ala 355 360 365

Gly Pro Cys Cys Thr Pro Thr Lys Met Ser Pro Ile Asn Met Leu Tyr 370 375 380

Phe Asn Asp Lys Gln Gln Ile Ile Tyr Gly Lys Ile Pro Gly Met Val 385 390 395 400

Val Asp Arg Cys Gly Cys Ser 405

<210> 296

<211> 405

<212> PRT

<213> Mus musculus

<400> 296

Met Val Leu Ala Ala Pro Leu Leu Leu Gly Phe Leu Leu Leu Ala Leu 1 5 10 15

Glu Leu Arg Pro Arg Gly Glu Ala Ala Glu Gly Pro Ala Ala Ala Ala 20 25 30

Ala Ala Ala Ala Ala Ala Gly Val Gly Glu Arg Ser Ser Arg
35 40 45

Pro Ala Pro Ser Ala Pro Pro Glu Pro Asp Gly Cys Pro Val Cys Val
50 55 60

Trp Arg Gln His Ser Arg Glu Leu Arg Leu Glu Ser Ile Lys Ser Gln 65 70 75 80

Ile Leu Ser Lys Leu Arg Leu Lys Glu Ala Pro Asn Ile Ser Arg Glu 85 90 95

Val Val Lys Gln Leu Leu Pro Lys Ala Pro Pro Leu Gln Gln Ile Leu 100 105 110

| Asp | Leu | His 115 | Asp | Phe | Gln | Gly | Asp 120 | Ala | Leu | Gln | Pro | Glu 125 | Asp | Phe | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu | Glu 130 | Asp | Glu | Tyr | His | Ala 135 | Thr | Thr | Glu | Thr | Val 140 | Ile | Ser | Met | Ala |
| Gln 145 | Glu | Thr | Asp | Pro | Ala 150 | Val | Gln | Thr | Asp | Gly 155 | Ser | Pro | Leu | Cys | Cys 160 |
| His | Phe | His | Phe | Ser 165 | Pro | Lys | Val | Met | Phe 170 | Thr | Lys | Val | Leu | Lys 175 | Ala |
| Gln | Leu | Trp | Val 180 | Tyr | Leu | Arg | Pro | Val 185 | Pro | Arg | Pro | Ala | Thr 190 | Val | Tyr |
| Leu | Gln | Ile 195 | Leu | Arg | Leu | Lys | Pro 200 | Leu | Thr | Gly | Glu | Gly 205 | Thr | Ala | Gly |
| Gly | Gly 210 | Gly | Gly | Gly | Arg | Arg 215 | His | Ile | Arg | Ile | Arg 220 | Ser | Leu | Lys | Ile |
| Glu 225 | Leu | His | Ser | Arg | Ser 230 | Gly | His | Trp | Gln | Ser 235 | Ile | Asp | Phe | Lys | Gln 240 |
| Val | Leu | His | Ser | Trp 245 | Phe | Arg | Gln | Pro | Gln 250 | Ser | Asn | Trp | Gly | Ile 255 | Glu |
| Ile | Asn | Ala | Phe 260 | Asp | Pro | Ser | Gly | Thr 265 | Asp | Leu | Ala | Val | Thr 270 | Ser | Leu |
| Gly | Pro | Gly 275 | Ala | Glu | Gly | Leu | His 280 | Pro | Phe | Met | Glu | Leu 285 | Arg | Val | Leu |
| Glu | Asn 290 | Thr | Lys | Arg | Ser | Arg 295 | Arg | Asn | Leu | Gly | Leu 300 | Asp | Cys | Asp | Glu |
| His 305 | Ser | Ser | Glu | Ser | Arg 310 | Cys | Cys | Arg | Tyr | Pro 315 | Leu | Thr | Val | Asp | Phe 320 |
| Glu | Ala | Phe | Gly | Trp 325 | Asp | Trp | Ile | Ile | Ala 330 | Pro | Lys | Arg | Tyr | Lys 335 | Ala |
| Asn | Tyr | Cys | Ser 340 | Gly | Gln | Cys | Glu | Tyr 345 | Met | Phe | Met | Gln | Lys 350 | Tyr | Pro |
| His | Thr | His 355 | Leu | Val | Gln | Gln | Ala 360 | Asn | Pro | Arg | Gly | Ser 365 | Ala | Gly | Pro |

Cys Cys Thr Pro Thr Lys Met Ser Pro Ile Asn Met Leu Tyr Phe Asn 370 375 380

Asp Lys Gln Gln Ile Ile Tyr Gly Lys Ile Pro Gly Met Val Val Asp 385 390 395 400

Arg Cys Gly Cys Ser 405

<210> 297

<211> 405

<212> PRT

<213> Mus musculus

<400> 297

Met Val Leu Ala Ala Pro Leu Leu Gly Phe Leu Leu Leu Ala Leu 1 5 10 15

Glu Leu Arg Pro Arg Gly Glu Ala Ala Glu Gly Pro Ala Ala Ala Ala 20 25 30

Ala Ala Ala Ala Ala Ala Gly Val Gly Glu Arg Ser Ser Arg
35 40 45

Pro Ala Pro Ser Ala Pro Pro Glu Pro Asp Gly Cys Pro Val Cys Val 50 55 60

Trp Arg Gln His Ser Arg Glu Leu Arg Leu Glu Ser Ile Lys Ser Gln 65 70 75 80

Ile Leu Ser Lys Leu Arg Leu Lys Glu Ala Pro Asn Ile Ser Arg Glu 85 90 95

Val Val Lys Gln Leu Leu Pro Lys Ala Pro Pro Leu Gln Gln Ile Leu 100 105 110

Asp Leu His Asp Phe Gln Gly Asp Ala Leu Gln Pro Glu Asp Phe Leu
115 120 125

Glu Glu Asp Glu Tyr His Ala Thr Thr Glu Thr Val Ile Ser Met Ala 130 135 140

Gln Glu Thr Asp Pro Ala Val Gln Thr Asp Gly Ser Pro Leu Cys Cys 145 150 155 160

His Phe His Phe Ser Pro Lys Val Met Phe Asn Lys Val Leu Lys Ala 165 170 175

Gln Leu Trp Val Tyr Leu Arg Pro Val Pro Arg Pro Ala Thr Val Tyr Leu Gln Ile Leu Arg Leu Lys Pro Leu Thr Gly Glu Gly Thr Ala Gly Gly Gly Gly Gly Arg Arg His Ile Arg Ile Arg Ser Leu Lys Ile Glu Leu His Ser Arg Ser Gly His Trp Gln Ser Ile Asp Phe Lys Gln Val Leu His Ser Trp Phe Arg Gln Pro Gln Ser Asn Trp Gly Ile Glu Ile Asn Ala Phe Asp Pro Ser Gly Thr Asp Leu Ala Val Thr Ser Leu Gly Pro Gly Ala Glu Gly Leu His Pro Phe Met Glu Leu Arg Val Leu Glu Asn Thr Lys Arg Ser Arg Arg Asn Leu Gly Leu Asp Cys Asp Glu His Ser Ser Glu Ser Arg Cys Cys Arg Tyr Pro Leu Thr Val Asp Phe Glu Ala Phe Gly Trp Asp Trp Ile Ile Ala Pro Lys Arg Tyr Lys Ala Asn Tyr Cys Ser Gly Gln Cys Glu Tyr Met Phe Met Gln Lys Tyr Pro His Thr His Leu Val Gln Gln Ala Asn Pro Arg Gly Ser Ala Gly Pro Cys Cys Thr Pro Thr Lys Met Ser Pro Ile Asn Met Leu Tyr Phe Asn Asp Lys Gln Gln Ile Ile Tyr Gly Lys Ile Pro Gly Met Val Val Asp Arg Cys Gly Cys Ser

| <211 | L> 34 | 15 | | | |
|-------|-------|-------|-------|-------|------|
| <212 | 2> PE | RT | | | |
| <213 | 3> Ra | attus | s noi | cvegi | icus |
| | | | | | |
| < 400 |)> 29 | 98 | | | |
| Pro | Glu | Pro | Asp | Gly | Cys |
| 1 | | | | 5 | |
| | | | | | |
| Arg | Val | Arg | Leu | Gly | Ser |

- Pro Glu Pro Asp Gly Cys Pro Val Cys Val Trp Arg Gln His Ser Arg 1 5 10 15
- Arg Val Arg Leu Gly Ser Ile Lys Ser Gln Ile Leu Ser Lys Leu Arg 20 25 30
- Leu Lys Glu Ala Pro Asn Ile Ser Arg Glu Val Val Lys Gln Leu Leu 35 40 45
- Pro Lys Ala Pro Pro Leu Gln Gln Ile Leu Asp Leu His Asp Phe Gln 50 55 60
- Gly Asp Ala Leu Gln Pro Glu Asp Phe Leu Glu Glu Asp Glu Tyr His 65 70 75 80
- Ala Thr Thr Glu Thr Val Ile Ser Met Ala Gln Glu Thr Asp Pro Ala 85 90 95
- Val Gln Thr Asp Gly Ser Pro Leu Cys Cys His Phe His Phe Ser Pro 100 105 110
- Lys Val Met Phe Thr Lys Val Leu Lys Ala Gln Leu Trp Val Tyr Leu
 115 120 125
- Arg Pro Val Pro Arg Pro Ala Thr Val Tyr Leu Gln Ile Leu Arg Leu 130 135 140
- Lys Pro Leu Thr Gly Glu Gly Thr Ala Gly Gly Gly Gly Gly Gly Arg
 145 150 155 160
- Arg His Ile Arg Ile Arg Ser Leu Lys Ile Glu Leu His Ser Arg Ser 165 170 175
- Gly His Trp Gln Ser Ile Asp Phe Lys Gln Val Leu His Ser Trp Phe 180 185 190
- Arg Gln Pro Gln Ser Asn Trp Gly Ile Glu Ile Asn Ala Phe Asp Pro 195 200 205
- Ser Gly Thr Asp Leu Ala Val Thr Ser Leu Gly Pro Gly Ala Glu Gly 210 215 220
- Cys His Pro Phe Met Glu Leu Arg Val Leu Glu Asn Thr Lys Arg Ser

| 225 | 230 | 235 | 240 |
|----------------------------|--------------------|--------------------------|------------------------|
| Arg Arg Asn Leu Gly 245 | | Asp Glu His Ser S 250 | Ser Glu Ser Arg 255 |
| Cys Cys Arg Tyr Pro 260 | Leu Thr Val | Asp Phe Glu Ala S 265 | Ser Gly Trp Asp 270 |
| Trp Ile Ile Ala Pro 275 | Lys Arg Tyr 280 | - | Cys Ser Gly Gln 285 |
| Cys Glu Tyr Met Phe 290 | Met Gln Lys 295 | Tyr Pro His Thr F | His Leu Val Gln |
| Gln Ala Asn Pro Arg 305 | Gly Ser Ala 310 | Gly Pro Cys Cys 1 | Thr Pro Thr Lys 320 |
| Met Ser Pro Ile Asn 325 | _ | Phe Asn Asp Lys 0 | Gln Gln Ile Ile 335 |
| Tyr Gly Lys Ile Pro | Gly Met Val | Val 345 | |

<210> 299 <211> 95 <212> PRT

<213> Homo sapiens

<400> 299

Cys Cys Arg Tyr Pro Leu Thr Val Asp Phe Glu Ala Phe Gly Trp Asp 1 5 10 15

Trp Ile Ile Ala Pro Lys Arg Tyr Lys Ala Asn Tyr Cys Ser Gly Gln 20 25 30

Gln Ala Asn Pro Arg Gly Ser Ala Gly Pro Cys Cys Thr Pro Thr Lys
50 55 60

Met Ser Pro Ile Asn Met Leu Tyr Phe Asn Asp Lys Gln Gln Ile Ile 65 70 75 80

Tyr Gly Lys Ile Pro Gly Met Val Val Asp Arg Cys Gly Cys Ser 85 90 95

<211> 102 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Transforming growth factor beta like domain sequence <400> 300 Cys Arg Arg His Asp Leu Tyr Val Asp Phe Lys Asp Leu Gly Trp Asp 5 10 Asp Trp Ile Ile Ala Pro Lys Gly Tyr Asn Ala Tyr Tyr Cys Glu Gly 20 25 30 Glu Cys Pro Phe Pro Leu Ser Glu Arg Leu Asn Ala Thr Asn His Ala 35 40 45 Ile Val Gln Ser Leu Val His Ala Leu Asp Pro Gly Ala Val Pro Lys 50 55 Pro Cys Cys Val Pro Thr Lys Leu Ser Pro Leu Ser Met Leu Tyr Tyr 65 70 75 80 Asp Asp Asp Gly Asn Val Val Leu Arg Asn Tyr Pro Asn Met Val Val 85 90 Glu Glu Cys Gly Cys Arg 100 <210> 301 <211> 102 <212> PRT <213> Homo sapiens <400> 301 Cys Arg Leu Arg Ser Leu Tyr Val Asp Phe Arg Asp Leu Gly Trp Gly 1 5 10 15 Asp Trp Ile Ile Ala Pro Glu Gly Tyr Ile Ala Asn Tyr Cys Ser Gly 20 25 Ser Cys Pro Phe Pro Leu Arg Asp Asp Leu Asn Leu Ser Asn His Ala

<210> 300

40

45

Ile Leu Gln Thr Leu Val Arg Leu Arg Asn Pro Arg Ala Val Pro Gln 50 55 60

Pro Cys Cys Val Pro Thr Lys Leu Ser Pro Leu Ser Met Leu Tyr Leu 65 70 75 80

Asp Asp Asn Ser Asn Val Val Leu Arg Leu Tyr Pro Asn Met Ser Val 85 90 95

Lys Glu Cys Gly Cys Arg 100

<210> 302

<211> 105

<212> PRT

<213> Homo sapiens

<400> 302

Cys Pro Val Cys Val Trp Arg Gln His Ser Arg Glu Leu Arg Leu Glu
1 5 10 15

Ser Ile Lys Ser Gln Ile Leu Ser Lys Leu Arg Leu Lys Glu Ala Pro 20 25 30

Asn Ile Ser Arg Glu Val Val Lys Gln Leu Leu Pro Lys Ala Pro Pro 35 40 45

Leu Gln Gln Ile Leu Asp Leu His Asp Phe Gln Gly Asp Ala Leu Gln 50 55 60

Pro Glu Asp Phe Leu Glu Glu Asp Glu Tyr His Ala Thr Thr Glu Thr 65 70 75 80

Val Ile Ser Met Ala Gln Glu Thr Asp Pro Ala Val Gln Thr Asp Gly
85 90 95

Ser Pro Leu Cys Cys His Phe His Phe 100 105

<210> 303

<211> 105

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: TGF-beta

propeptide domain sequence

Cys Arg Pro Leu Asp Leu Arg Arg Ser Gln Lys Gln Asp Arg Leu Glu Ala Ile Glu Gly Gln Ile Leu Ser Lys Leu Gly Leu Arg Arg Pro Arg Pro Ser Lys Glu Pro Met Val Val Pro Glu Tyr Met Leu Asp Leu Tyr Asn Ala Leu Ser Glu Leu Glu Glu Gly Lys Val Gly Arg Val Pro Glu Ile Ser Asp Tyr Asp Gly Arg Glu Ala Gly Arg Ala Asn Thr Ile Arg Ser Phe Ser His Leu Glu Ser Asp Phe Glu Glu Ser Thr Pro Glu Ser His Arg Lys Arg Phe Arg Phe <210> 304 <211> 404 <212> PRT <213> Homo sapiens Met Ser Val Lys Pro Ser Trp Gly Pro Gly Pro Ser Glu Gly Val Thr Ala Val Pro Thr Ser Asp Leu Gly Glu Ile His Asn Trp Thr Glu Leu Leu Asp Leu Phe Asn His Thr Leu Ser Glu Cys His Val Glu Leu Ser Gln Ser Thr Lys Arg Val Val Leu Phe Ala Leu Tyr Leu Ala Met Phe Val Val Gly Leu Val Glu Asn Leu Leu Val Ile Cys Val Asn Trp Arg Gly Ser Gly Arg Ala Gly Leu Met Asn Leu Tyr Ile Leu Asn Met Ala

| Ile | Ala | Asp | Leu 100 | Gly | Ile | Val | Leu | Ser 105 | Leu | Pro | Val | Trp | Met 110 | Leu | Glu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val | Thr | Leu 115 | Asp | Tyr | Thr | Trp | Leu 120 | Trp | Gly | Ser | Phe | Ser 125 | Cys | Arg | Phe |
| Thr | His 130 | Tyr | Phe | Tyr | Phe | Val 135 | Asn | Met | Tyr | Ser | Ser 140 | Ile | Phe | Phe | Leu |
| Val 145 | Cys | Leu | Ser | Val | Asp 150 | Arg | Tyr | Val | Thr | Leu 155 | Thr | Ser | Ala | Ser | Pro 160 |
| Ser | Trp | Gln | Arg | Tyr 165 | Gln | His | Arg | Val | Arg 170 | Arg | Ala | Met | Cys | Ala 175 | Gly |
| Ile | Trp | Val | Leu 180 | Ser | Ala | Ile | Ile | Pro 185 | Leu | Pro | Glu | Val | Val 190 | His | Ile |
| Gln | Leu | Val 195 | Glu | Gly | Pro | Glu | Pro 200 | Met | Cys | Leu | Phe | Met 205 | Ala | Pro | Phe |
| Glu | Thr 210 | Tyr | Ser | Thr | Trp | Ala 215 | Leu | Ala | Val | Ala | Leu 220 | Ser | Thr | Thr | Ile |
| Leu 225 | Gly | Phe | Leu | Leu | Pro 230 | Phe | Pro | Leu | Ile | Thr 235 | Val | Phe | Asn | Val | Leu 240 |
| Thr | Ala | Cys | Arg | Leu 245 | Arg | Gln | Pro | Gly | Gln 250 | Pro | Lys | Ser | Arg | Arg 255 | His |
| Cys | Leu | Leu | Leu 260 | Cys | Ala | Tyr | Val | Ala 265 | Val | Phe | Val | Met | Cys 270 | Trp | Leu |
| Pro | Tyr | His 275 | Val | Thr | Leu | Leu | Leu 280 | Leu | Thr | Leu | His | Gly 285 | Thr | His | Ile |
| Ser | Leu 290 | His | Cys | His | Leu | Val 295 | His | Leu | Leu | Tyr | Phe 300 | Phe | Tyr | Asp | Val |
| Ile 305 | Asp | Cys | Phe | Ser | Met 310 | Leu | His | Cys | Val | Ile 315 | Asn | Pro | Ile | Leu | Tyr 320 |
| Asn | Phe | Leu | Ser | Pro 325 | His | Phe | Arg | Gly | Arg 330 | Leu | Leu | Asn | Ala | Val 335 | Val |
| His | Tyr | Leu | Pro 340 | Lys | Asp | Gln | Thr | Lys 345 | Ala | Gly | Thr | Cys | Ala 350 | Ser | Ser |

Ser Ser Cys Ser Thr Gln His Ser Ile Ile Ile Thr Lys Gly Asp Ser 355 360 365

Gln Pro Ala Ala Ala Ala Pro His Pro Glu Pro Ser Leu Ser Phe Gln 370 380

Ala His His Leu Leu Pro Asn Thr Ser Pro Ile Ser Pro Thr Gln Pro 385 390 395 400

Leu Thr Pro Ser

<210> 305

<211> 395

<212> PRT

<213> Mus musculus

<400> 305

Met Ser Val Ile Pro Ser Pro Arg Pro Val Ser Thr Leu Glu Pro Asp 1 5 10 15

Asn Asp Phe Arg Asp Ile His Asn Trp Thr Glu Leu Leu His Leu Phe
20 25 30

Asn Gln Thr Phe Thr Asp Cys His Ile Glu Phe Asn Glu Asn Thr Lys
35 40 45

His Val Val Leu Phe Val Phe Tyr Leu Ala Ile Phe Val Val Gly Leu 50 55 60

Val Glu Asn Val Leu Val Ile Cys Val Asn Cys Arg Arg Ser Gly Arg 65 70 75 80

Val Gly Met Leu Asn Leu Tyr Ile Leu Asn Met Ala Ile Ala Asp Leu 85 90 95

Gly Ile Ile Leu Ser Leu Pro Val Trp Met Leu Glu Val Met Leu Glu 100 105 110

Tyr Thr Trp Leu Trp Gly Ser Phe Ser Cys Arg Phe Ile His Tyr Phe
115 120 125

Tyr Leu Val Asn Met Tyr Ser Ser Ile Phe Phe Leu Thr Cys Leu Ser 130 135 140

Ile Asp Arg Tyr Val Thr Leu Thr Asn Thr Ser Pro Ser Trp Gln Arg

| 145 | 150 | | 155 | 160 |
|------------------------|--------------------|--------------------|------------------------|----------------|
| His Gln His Arg | Ile Arg Arg 165 | Ala Val Cys 170 | Ala Gly Val Trp | Val Leu 175 |
| Ser Ala Ile Ile 180 | Pro Leu Pro | Glu Val Val 185 | His Ile Gln Leu 190 | Leu Asp |
| Gly Ser Glu Pro 195 | Met Cys Leu | Phe Leu Ala 200 | Pro Phe Glu Thr 205 | Tyr Ser |
| Ala Trp Ala Leu 210 | Ala Val Ala 215 | Leu Ser Ala | Thr Ile Leu Gly 220 | Phe Leu |
| Leu Pro Phe Leu 225 | Leu Ile Ala 230 | Val Phe Asn | Ile Leu Thr Ala 235 | Cys Arg 240 |
| Leu Arg Arg Gln | Arg Gln Thr 245 | Glu Ser Arg 250 | Arg His Cys Leu | Leu Met 255 |
| Trp Ala Tyr Ile 260 | Val Val Phe | Ala Ile Cys 265 | Trp Leu Pro Tyr 270 | Gln Val |
| Thr Met Leu Leu 275 | Leu Thr Leu | His Gly Thr 280 | His Ile Phe Leu 285 | His Cys |
| His Leu Val Asn 290 | Leu Leu Tyr 295 | Phe Phe Tyr | Glu Ile Ile Asp 300 | Cys Phe |
| Ser Met Leu His 305 | Cys Val Ala 310 | Asn Pro Ile | Leu Tyr Asn Phe 315 | Leu Ser 320 |
| Pro Ser Phe Arg | Gly Arg Leu 325 | Leu Ser Leu 330 | Val Val Arg Tyr | Leu Pro 335 |
| Lys Glu Gln Ala 340 | Arg Ala Ala | Gly Gly Arg 345 | Ala Ser Ser Ser 350 | Ser Ser |
| Thr Gln His Ser 355 | Ile Ile Ile | Thr Lys Glu 360 | Gly Ser Leu Pro 365 | Leu Gln |
| Arg Ile Ser Thr 370 | Pro Thr Pro 375 | Ser Glu Thr | Phe Arg Arg Pro 380 | Leu Arg |
| Leu Gln Thr Pro 385 | His Leu His 390 | Ser Ala Ile | Leu 395 | |

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<210> 306
<211> 398
<212> PRT
<213> Rattus norvegicus
<400> 306
Met Ser Val Ile Pro Ser
1 5
```

Met Ser Val Ile Pro Ser Ser Arg Pro Val Ser Thr Leu Ala Pro Asp 1 5 10 15

Asn Asp Phe Arg Glu Ile His Asn Trp Thr Glu Leu Leu His Leu Phe 20 25 30

Asn Gln Thr Phe Ser Asp Cys Arg Met Glu Leu Asn Glu Asn Thr Lys 35 40 45

Gln Val Val Leu Phe Val Phe Tyr Leu Ala Ile Phe Val Val Gly Leu 50 55 60

Val Glu Asn Val Leu Val Ile Cys Val Asn Cys Arg Arg Ser Gly Arg 65 70 75 80

Val Gly Met Leu Asn Leu Tyr Ile Leu Asn Met Ala Val Ala Asp Leu 85 90 95

Gly Ile Ile Leu Ser Leu Pro Val Trp Met Leu Glu Val Met Leu Glu
100 105 110

Tyr Thr Trp Leu Trp Gly Ser Phe Ser Cys Arg Phe Ile His Tyr Phe
115 120 125

Tyr Leu Ala Asn Met Tyr Ser Ser Ile Phe Phe Leu Thr Cys Leu Ser 130 135 140

Ile Asp Arg Tyr Val Thr Leu Thr Asn Thr Ser Pro Ser Trp Gln Arg
145 150 155 160

His Gln His Arg Ile Arg Arg Ala Val Cys Ala Gly Val Trp Val Leu 165 170 175

Ser Ala Ile Ile Pro Leu Pro Glu Val Val His Ile Gln Leu Leu Asp 180 185 190

Gly Ser Glu Pro Met Cys Leu Phe Leu Ala Pro Phe Glu Thr Tyr Ser 195 200 205

Ala Trp Ala Leu Ala Val Ala Leu Ser Ala Thr Ile Leu Gly Phe Leu 210 215 220

| 225 | PIO | Pne | Pro | Leu | 230 | АІА | vai | rne | Asn | 235 | Leu | ser | АТА | cys | 240 |
|--------------|----------------------------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Arg | Arg | Gln | Gly 245 | Gln | Thr | Glu | Ser | Arg 250 | Arg | His | Cys | Leu | Leu 255 | Met |
| Trp | Ala | Tyr | Ile 260 | Val | Val | Phe | Ala | Ile 265 | Cys | Trp | Leu | Pro | Tyr 270 | His | Val |
| Thr | Met | Leu 275 | Leu | Leu | Thr | Leu | His 280 | Thr | Thr | His | Ile | Phe 285 | Leu | His | Cys |
| Asn | Leu 290 | Val | Asn | Phe | Leu | Tyr 295 | Phe | Phe | Tyr | Glu | Ile 300 | Thr | Asp | Cys | Phe |
| Ser 305 | Met | Leu | His | Cys | Val 310 | Ala | Asn | Pro | Ile | Leu 315 | Tyr | Asn | Phe | Leu | Ser 320 |
| Pro | Ser | Phe | Arg | Gly 325 | Arg | Leu | Leu | Ser | Leu 330 | Val | Val | Arg | Tyr | Leu 335 | Pro |
| Lys | Glu | Gln | Ala 340 | Arg | Ala | Ala | Gly | Gly 345 | Arg | Ala | Ser | Ser | Ser 350 | Ser | Ser |
| Thr | Gln | His 355 | Ser | Ile | Ile | Ile | Thr 360 | Lys | Glu | Gly | Ser | Leu 365 | Leu | Ala | Ala |
| Ala | Asp 370 | Leu | His | Thr | His | Ala 375 | Ile | Arg | Asn | Val | Gln 380 | Ala | Ser | Ser | Leu |
| Pro 385 | Pro | Asn | Thr | Ser | Pro 390 | Thr | Leu | Cys | Asn | Ser 395 | Ile | Ala | Ser | | |
| <211 <212 |)> 3(l> 39 2> PI 3> Ra | 95 RT | s noi | rveg: | icus | | | | | | | | | | |
| |)> 3(Ser | | Tle | Pro | Ser | Ser | Glu | Ala | Val | Ser | Thr | Len | Ala | Pro | Asp |
| 1 | | ·uı | | 5 | JU1 | | Jiu | | 10 | | -111 | u | | 15 | |
| Asn | Asp | Phe | Arg 20 | Glu | Ile | His | Asn | Trp 25 | Thr | Glu | Leu | Leu | His 30 | Leu | Phe |
| Asn | Gln | Thr 35 | Phe | Ser | Asp | Cys | His 40 | Met | Glu | Leu | Asn | Glu 45 | Asn | Thr | Lys |

Leu Pro Phe Pro Leu Ile Ala Val Phe Asn Ile Leu Ser Ala Cys Arg

| Gln | Val 50 | Val | Leu | Phe | Val | Phe 55 | Tyr | Leu | Ala | Ile | Phe 60 | Val | Val | Gly | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val 65 | Glu | Asn | Val | Leu | Val 70 | Ile | Cys | Val | Asn | Cys 75 | Arg | Arg | Ser | Gly | Arg 80 |
| Val | Gly | Met | Leu | Asn 85 | Leu | Tyr | Ile | Leu | Asn 90 | Met | Ala | Val | Ala | Asp 95 | Leu |
| Gly | Ile | Ile | Leu 100 | Ser | Leu | Pro | Val | Trp 105 | Met | Leu | Glu | Val | Met 110 | Leu | Glu |
| Tyr | Thr | Trp 115 | Leu | Trp | Gly | Ser | Phe 120 | Ser | Cys | Arg | Phe | Ile 125 | His | Tyr | Phe |
| Tyr | Leu 130 | Ala | Asn | Met | Tyr | Ser 135 | Ser | Ile | Phe | Phe | Leu 140 | Thr | Cys | Leu | Ser |
| Ile 145 | Asp | Arg | Tyr | Val | Thr 150 | Leu | Thr | Asn | Thr | Ser 155 | Pro | Ser | Trp | Gln | Arg 160 |
| His | Gln | His | Arg | Ile 165 | Arg | Arg | Ala | Val | Cys 170 | Ala | Gly | Val | Trp | Val 175 | Leu |
| Ser | Ala | Ile | Ile 180 | Pro | Leu | Pro | Glu | Val 185 | Val | His | Ile | Gln | Leu 190 | Leu | Asp |
| Gly | Ser | Glu 195 | Pro | Met | Cys | Leu | Phe 200 | Leu | Ala | Pro | Phe | Glu 205 | Thr | Tyr | Ser |
| Ala | Trp 210 | Ala | Leu | Ala | Val | Ala 215 | Leu | Ser | Ala | Thr | Ile 220 | Leu | Gly | Phe | Leu |
| Leu 225 | Pro | Phe | Pro | Leu | Ile 230 | Ala | Val | Phe | Asn | Ile 235 | Leu | Ser | Ala | Cys | Arg 240 |
| Leu | Arg | Arg | Gln | Gly 245 | Gln | Thr | Glu | Ser | Arg 250 | Arg | His | Суѕ | Leu | Leu 255 | Met |
| Trp | Ala | Tyr | Ile 260 | Ala | Val | Phe | Val | Ile 265 | Cys | Trp | Leu | Pro | Tyr 270 | His | Val |
| Thr | Met | Leu 275 | Leu | Leu | Thr | Leu | His 280 | Thr | Thr | His | Ile | Phe 285 | Leu | His | Cys |
| Asn | Leu 290 | Val | Asn | Phe | Leu | Tyr 295 | Phe | Phe | Tyr | Glu | Ile 300 | Ile | Asp | Cys | Phe |

Ser Met Leu His Cys Val Ala Asn Pro Ile Leu Tyr Asn Phe Leu Ser 305 310 315 320

Pro Ser Phe Arg Gly Arg Leu Leu Ser Leu Val Val Arg Tyr Leu Pro 325 330 335

Lys Glu Gln Ala Arg Ala Ala Gly Gly Arg Ala Ser Ser Ser Ser Ser 340 345 350

Thr Gln His Ser Ile Ile Ile Thr Lys Glu Gly Ser Leu Pro Leu Gln 355 360 365

Arg Ile Cys Thr Pro Thr Pro Ser Glu Thr Cys Arg Pro Pro Leu Cys 370 380

Leu Arg Thr Pro His Leu His Ser Ala Ile Pro 385 390 395

<210> 308

<212> PRT

<213> Rattus norvegicus

<400> 308

<210> 309

<211> 75

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7tm_1: domain
 sequence

<400> 309

Gly Asn Leu Leu Val Ile Leu Val Ile Leu Arg Thr Lys Lys Leu Arg 1 5 10 15

Thr Pro Thr Asn Ile Phe Ile Leu Asn Leu Ala Val Ala Asp Leu Leu 20 25 30

Phe Leu Leu Thr Leu Pro Pro Trp Ala Leu Tyr Tyr Leu Val Gly Gly 35 40 45

Ser Glu Asp Trp Pro Phe Gly Ser Ala Leu Cys Lys Leu Val Thr Ala

50 55 60

Leu Asp Val Val Asn Met Tyr Ala Ser Ile Leu 65 70 75

<210> 310

<211> 73

<212> PRT

<213> Homo sapiens

<400> 310

Glu Asn Leu Leu Val Ile Cys Val Asn Trp Arg Gly Ser Gly Arg Ala 1 5 10 15

Gly Leu Met Asn Leu Tyr Ile Leu Asn Met Ala Ile Ala Asp Leu Gly
20 25 30

Ile Val Leu Ser Leu Pro Val Trp Met Leu Glu Val Thr Leu Asp Tyr 35 40 45

Thr Trp Leu Trp Gly Ser Phe Ser Cys Arg Phe Thr His Tyr Phe Tyr 50 55 60

Phe Val Asn Met Tyr Ser Ser Ile Phe 65 70

<210> 311

<211> 87

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7tm_1: domain
 sequence

<400> 311

Phe Leu Leu Pro Leu Leu Val Ile Leu Val Cys Tyr Thr Arg Ile Leu

1 5 10 15

Arg Thr Leu Arg Lys Ala Ala Lys Thr Leu Leu Val Val Val Val Val Val 20 25 30

Phe Val Leu Cys Trp Leu Pro Tyr Phe Ile Val Leu Leu Leu Asp Thr 35 40 45

Leu Cys Leu Ser Ile Ile Met Ser Ser Thr Cys Glu Leu Glu Arg Val

50 55 60

Leu Pro Thr Ala Leu Leu Val Thr Leu Trp Leu Ala Tyr Val Asn Ser 65 70 75 80

Cys Leu Asn Pro Ile Ile Tyr 85

<210> 312

<211> 94

<212> PRT

<213> Homo sapiens

<400> 312

Phe Leu Leu Pro Phe Pro Leu Ile Thr Val Phe Asn Val Leu Thr Ala 1 5 10 15

Cys Arg Leu Arg Gln Pro Gly Gln Pro Lys Ser Arg Arg His Cys Leu 20 25 30

Leu Leu Cys Ala Tyr Val Ala Val Phe Val Met Cys Trp Leu Pro Tyr 35 40 45

His Val Thr Leu Leu Leu Leu Thr Leu His Gly Thr His Ile Ser Leu 50 55 60

His Cys His Leu Val His Leu Leu Tyr Phe Phe Tyr Asp Val Ile Asp 65 70 75 80

Cys Phe Ser Met Leu His Cys Val Ile Asn Pro Ile Leu Tyr 85 90

<210> 313

<211> 254

<212> PRT

<213> Homo sapiens

<400> 313

Gly Asn Leu Leu Val Ile Leu Val Ile Leu Arg Thr Lys Lys Leu Arg
1 5 10 15

Thr Pro Thr Asn Ile Phe Leu Leu Asn Leu Ala Val Ala Asp Leu Leu 20 25 30

Phe Leu Leu Thr Leu Pro Pro Trp Ala Leu Tyr Tyr Leu Val Gly Gly 35 40 45

Asp Trp Val Phe Gly Asp Ala Leu Cys Lys Leu Val Gly Ala Leu Phe 50 55 60 Val Val Asp Gly Tyr Ala Ser Ile Leu Leu Leu Thr Ala Ile Ser Ile

val Val Ash Gly Tyr Ala Ser lie Leu Leu Leu Thr Ala lie Ser lie
65 70 75 80

Asp Arg Tyr Leu Ala Ile Val His Pro Leu Arg Tyr Arg Arg Ile Arg 85 90 95

Thr Pro Arg Arg Ala Lys Val Leu Ile Leu Leu Val Trp Val Leu Ala 100 105 110

Leu Leu Ser Leu Pro Pro Leu Leu Phe Ser Trp Leu Arg Thr Val 115 120 125

Glu Glu Gly Asn Thr Thr Val Cys Leu Ile Asp Phe Pro Glu Glu Ser 130 135 140

Val Lys Arg Ser Tyr Val Leu Leu Ser Thr Leu Val Gly Phe Val Leu 145 150 155 160

Pro Leu Leu Val Ile Leu Val Cys Tyr Thr Arg Ile Leu Arg Thr Leu 165 170 175

Arg Lys Arg Ala Arg Ser Gln Arg Ser Leu Lys Arg Arg Ser Ser Ser 180 185 190

Glu Arg Lys Ala Ala Lys Met Leu Leu Val Val Val Val Phe Val
195 200 205

Leu Cys Trp Leu Pro Tyr His Ile Val Leu Leu Leu Asp Ser Leu Cys 210 215 220

Leu Leu Ser Ile Trp Arg Val Leu Pro Thr Ala Leu Leu Ile Thr Leu 225 230 235 240

Trp Leu Ala Tyr Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr
245 250

<210> 314

<211> 254

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: 7tm_1: domain

sequence

| < 400 |)> 31 | L 4 | | | | | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly 1 | Asn | Leu | Leu | Val 5 | Ile | Leu | Val | Ile | Leu 10 | Arg | Thr | Lys | Lys | Leu 15 | Arg |
| Thr | Pro | Thr | Asn 20 | Ile | Phe | Leu | Leu | Asn 25 | Leu | Ala | Val | Ala | Asp 30 | Leu | Leu |
| Phe | Leu | Leu 35 | Thr | Leu | Pro | Pro | Trp 40 | Ala | Leu | Tyr | Tyr | Leu 45 | Val | Gly | Gly |
| Asp | Trp 50 | Val | Phe | Gly | Asp | Ala 55 | Leu | Cys | Lys | Leu | Val 60 | Gly | Ala | Leu | Phe |
| Val 65 | Val | Asn | Gly | Tyr | Ala 70 | Ser | Ile | Leu | Leu | Leu 75 | Thr | Ala | Ile | Ser | Ile 80 |
| Asp | Arg | Tyr | Leu | Ala 85 | Ile | Val | His | Pro | Leu 90 | Arg | Tyr | Arg | Arg | Ile 95 | Arg |
| Thr | Pro | Arg | Arg 100 | Ala | Lys | Val | Leu | Ile 105 | Leu | Leu | Val | Trp | Val 110 | Leu | Ala |
| Leu | Leu | Leu 115 | Ser | Leu | Pro | Pro | Leu 120 | Leu | Phe | Ser | Trp | Leu 125 | Arg | Thr | Val |
| Glu | Glu 130 | Gly | Asn | Thr | Thr | Val 135 | Cys | Leu | Ile | Asp | Phe 140 | Pro | Glu | Glu | Ser |
| Val 145 | Lys | Arg | Ser | Tyr | Val 150 | Leu | Leu | Ser | Thr | Leu 155 | Val | Gly | Phe | Val | Leu 160 |
| Pro | Leu | Leu | Val | Ile 165 | Leu | Val | Cys | Tyr | Thr 170 | Arg | Ile | Leu | Arg | Thr 175 | Leu |
| Arg | Lys | Arg | Ala 180 | Arg | Ser | Gln | Arg | Ser 185 | Leu | Lys | Arg | Arg | Ser 190 | Ser | Ser |
| Glu | Arg | Lys 195 | Ala | Ala | Lys | Met | Leu 200 | Leu | Val | Val | Val | Val 205 | Val | Phe | Val |
| Leu | Cys 210 | Trp | Leu | Pro | Tyr | His 215 | Ile | Val | Leu | Leu | Leu 220 | Asp | Ser | Leu | Cys |
| Leu 225 | Leu | Ser | Ile | Trp | Arg 230 | Val | Leu | Pro | Thr | Ala 235 | Leu | Leu | Ile | Thr | Leu 240 |

Trp Leu Ala Tyr Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr 245 250

<210> 315

<211> 173

<212> PRT

<213> Homo sapiens

<400> 315

Met Ala Arg Met Asn Arg Pro Ala Pro Val Glu Val Thr Tyr Lys Asn 1 5 10 15

Met Arg Phe Leu Ile Thr His Asn Pro Thr Asn Ala Thr Leu Asn Lys 20 25 30

Phe Ile Glu Glu Leu Lys Lys Tyr Gly Val Thr Thr Ile Val Arg Val 35 40 45

Cys Glu Ala Thr Tyr Asp Thr Thr Leu Val Glu Lys Glu Gly Ile His 50 55 60

Val Leu Asp Trp Pro Phe Asp Asp Gly Ala Pro Pro Ser Asn Gln Ile 65 70 75 80

Val Asp Asp Trp Leu Ser Leu Val Lys Ile Lys Phe Arg Glu Glu Pro 85 90 95

Gly Cys Cys Ile Ala Val His Cys Val Ala Gly Leu Gly Arg Ala Pro 100 105 110

Val Leu Val Ala Leu Ala Leu Ile Glu Gly Gly Met Lys Tyr Glu Asp 115 120 125

Ala Val Gln Phe Ile Arg Gln Lys Arg Arg Gly Ala Phe Asn Ser Lys 130 135 140

Gln Leu Leu Tyr Leu Glu Lys Tyr Arg Pro Lys Met Arg Leu Arg Phe 145 150 155 160

Lys Asp Ser Asn Gly His Arg Asn Asn Cys Cys Ile Gln 165 170

<210> 316

<211> 173

<212> PRT

<213> Rattus norvegicus

| <400> 316 | | | | | | | | | | | | | | | |
|---|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-----------|------------|
| Met 1 | Ala | Arg | Met | Asn 5 | Arg | Pro | Ala | Pro | Val 10 | Glu | Val | Thr | Tyr | Lys 15 | Asn |
| Met | Arg | Phe | Leu 20 | Ile | Thr | His | Asn | Pro 25 | Thr | Asn | Ala | Thr | Leu 30 | Asn | Lys |
| Phe | Ile | Glu 35 | Glu | Leu | Lys | Lys | Tyr 40 | Gly | Val | Thr | Thr | Ile 45 | Val | Arg | Val |
| Cys | Glu 50 | Ala | Thr | Tyr | Asp | Thr 55 | Thr | Leu | Val | Glu | Lys 60 | Glu | Gly | Ile | His |
| Val 65 | Leu | Asp | Trp | Pro | Phe 70 | Asp | Asp | Gly | Ala | Pro 75 | Pro | Ser | Asn | Gln | Ile 80 |
| Val | Asp | Asp | Trp | Leu 85 | Ser | Leu | Val | Lys | Ile 90 | Lys | Phe | Arg | Glu | Glu 95 | Pro |
| Gly | Cys | Cys | Ile 100 | Ala | Val | His | Cys | Val 105 | Ala | Gly | Leu | Gly | Arg 110 | Ala | Pro |
| Val | Leu | Val 115 | Ala | Leu | Ala | Leu | Ile 120 | Glu | Gly | Gly | Met | Lys 125 | Tyr | Glu | Asp |
| Ala | Val 130 | Gln | Phe | Ile | Arg | Gln 135 | Lys | Arg | Arg | Gly | Ala 140 | Phe | Asn | Ser | Lys |
| Gln 145 | Leu | Leu | Tyr | Leu | Glu 150 | Lys | Tyr | Arg | Pro | Lys 155 | Met | Arg | Leu | Arg | Phe 160 |
| Lys | Asp | Ser | Asn | Gly 165 | His | Arg | Asn | Asn | Trp 170 | Cys | Ile | Gln | | | |
| <210> 317 <211> 167 <212> PRT <213> Homo sapiens | | | | | | | | | | | | | | | |

Glu Leu Lys Lys Tyr Gly Val Thr Thr Leu Val Arg Val Cys Asp Ala Thr Tyr Asp Lys Ala Pro Val Glu Lys Glu Gly Ile His Val Leu Asp Trp Pro Phe Asp Asp Gly Ala Pro Pro Pro Asn Gln Ile Val Asp Asp Trp Leu Asn Leu Leu Lys Thr Lys Phe Arg Glu Glu Pro Gly Cys Cys Val Ala Val His Cys Val Ala Gly Leu Gly Arg Ala Pro Val Leu Val Ala Leu Ala Leu Ile Glu Cys Gly Met Lys Tyr Glu Asp Ala Val Gln Phe Ile Arg Gln Lys Arg Arg Gly Ala Phe Asn Ser Lys Gln Leu Leu Tyr Leu Glu Lys Tyr Arg Pro Lys Met Arg Leu Arg Phe Arg Asp Thr Asn Gly His Cys Cys Val Gln <210> 318 <211> 167 <212> PRT <213> Homo sapiens <400> 318 Met Asn Arg Pro Ala Pro Val Glu Ile Ser Tyr Glu Asp Met Arg Phe Leu Ile Thr His Asn Pro Thr Asn Ala Thr Leu Asn Lys Phe Thr Glu Glu Leu Lys Lys Tyr Gly Val Thr Thr Leu Val Arg Val Cys Asp Ala Thr Tyr Asp Lys Ala Pro Val Glu Lys Glu Gly Ile His Val Leu Asp Trp Pro Phe Asp Asp Gly Ala Pro Pro Pro Asn Gln Ile Val Asp Asp

Trp Leu Asn Leu Leu Lys Thr Lys Phe Arg Glu Glu Pro Gly Cys Cys
85 90 95

Val Ala Val His Cys Val Ala Gly Leu Gly Arg Ala Pro Val Leu Val 100 105 110

Ala Leu Ala Leu Ile Glu Cys Gly Met Lys Tyr Glu Asp Ala Val Gln 115 120 125

Phe Ile Arg Gln Lys Arg Gly Ala Phe Asn Ser Lys Gln Leu Leu 130 135 140

Tyr Leu Glu Lys Tyr Arg Pro Lys Met Arg Leu Arg Phe Arg Asp Thr
145 150 155 160

Asn Gly His Cys Cys Val Gln 165

<210> 319

<211> 167

<212> PRT

<213> Mus musculus

<400> 319

Met Asn Arg Pro Ala Pro Val Glu Ile Ser Tyr Glu Asn Met Arg Phe 1 5 10 15

Leu Ile Thr His Asn Pro Thr Asn Ala Thr Leu Asn Lys Phe Thr Glu 20 25 30

Glu Leu Lys Lys Tyr Gly Val Thr Thr Leu Val Arg Val Cys Asp Ala
35 40 45

Thr Tyr Asp Lys Ala Pro Val Glu Lys Glu Gly Ile His Val Leu Asp 50 55 60

Trp Pro Phe Asp Asp Gly Ala Pro Pro Pro Asn Gln Ile Val Asp Asp 65 70 75 80

Trp Leu Asn Leu Leu Lys Thr Leu Phe Arg Glu Glu Pro Gly Cys Cys
85 90 95

Val Ala Val His Cys Val Ala Gly Ile Gly Arg Ala Pro Val Leu Val 100 105 110

Ala Leu Ala Leu Ile Glu Cys Gly Met Lys Tyr Glu Asp Ala Val Gln

115 120 125

Phe Ile Arg Gln Lys Arg Arg Gly Ala Phe Asn Ser Lys Gln Leu Leu 130 135 140

Tyr Leu Glu Lys Tyr Arg Pro Lys Met Arg Leu Arg Phe Arg Asp Thr 145 150 155 160

Asn Gly His Cys Cys Val Gln 165

<210> 320

<211> 130

<212> PRT

<213> Homo sapiens

<400> 320

Pro Ile Thr His Asn Pro Thr Asn Val Thr Leu Asn Lys Phe Ile Glu
1 5 10 15

Glu Leu Lys Lys Tyr Gly Ala Thr Thr Ile Val Arg Val Cys Glu Ala 20 25 30

Thr Tyr Asp Thr Thr Leu Val Glu Lys Glu Gly Ile His Val Leu Asn 35 40 45

Trp Pro Phe Gly Asp Gly Ala Pro Pro Ser Asn Gln Ile Val Ala Asp 50 55 60

Trp Leu His Phe Val Lys Ile Lys Phe Cys Glu Glu Pro Gly Cys Tyr 65 70 75 80

Ile Ala Val Asn Cys Ile Val Gly Leu Gly Lys Ala Pro Val Leu Val 85 90 95

Ala Leu Ala Ser Val Glu Gly Gly Met Lys His Glu Asp Ala Val Gln
100 105 110

Phe Ile Gly Gln Lys Arg Ser Gly Ala Phe Lys Ser Lys Gln Leu Leu 115 120 125

Tyr Leu

130

<210> 321

<211> 134

<212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Y_phosphatase domain sequence <400> 321 Ser Leu Thr Tyr Gly Asp Phe Thr Val Thr Cys Val Ser Val Glu Lys 5 10 Lys Lys Asp Asp Tyr Thr Val Arg Thr Leu Glu Leu Thr Asn Ser Gly 20 25 Asp Asp Glu Thr Arg Thr Val Lys His Tyr His Tyr Thr Gly Trp Pro 40 35 Asp His Gly Val Pro Glu Ser Pro Lys Ser Ile Leu Asp Leu Leu Arg 50 55 60 Lys Val Arg Lys Ser Lys Gly Thr Pro Asp Asp Gly Pro Ile Val Val 65 70 75 85 90

His Cys Ser Ala Gly Ile Gly Arg Thr Gly Thr Phe Ile Ala Ile Asp 95

80

Ile Leu Leu Gln Gln Leu Glu Lys Glu Gly Val Val Asp Val Phe Asp 100 105 110

Thr Val Lys Lys Leu Arg Ser Gln Arg Pro Gly Met Val Gln Thr Glu 115 120 125

Glu Gln Tyr Ile Phe Ile 130

<210> 322

<211> 90

<212> PRT

<213> Homo sapiens

<400> 322

His Val Leu Asn Trp Pro Phe Gly Asp Gly Ala Pro Pro Ser Asn Gln 5 15

Ile Val Ala Asp Trp Leu His Phe Val Lys Ile Lys Phe Cys Glu Glu 20 25 30

Pro Gly Cys Tyr Ile Ala Val Asn Cys Ile Val Gly Leu Gly Lys Ala 35 40 45

Pro Val Leu Val Ala Leu Ala Ser Val Glu Gly Gly Met Lys His Glu 50 55 60

Asp Ala Val Gln Phe Ile Gly Gln Lys Arg Ser Gly Ala Phe Lys Ser 65 70 75 80

Lys Gln Leu Leu Tyr Leu Glu Lys Tyr His 85 90

<210> 323

<211> 98

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: PTPc motif

<400> 323

His Tyr Thr Gly Trp Pro Asp His Gly Val Pro Glu Ser Pro Asp Ser 1 5 10 15

Ile Leu Glu Phe Leu Arg Ala Val Lys Lys Ser Leu Asn Lys Ser Ala 20 25 30

Asn Asn Gly Pro Val Val Val His Cys Ser Ala Gly Val Gly Arg Thr 35 40 45

Gly Thr Phe Val Ala Ile Asp Ile Leu Leu Gln Gln Leu Glu Ala Gly 50 55 . 60

Thr Gly Glu Val Asp Ile Phe Asp Ile Val Lys Glu Leu Arg Ser Gln 65 70 75 80

Arg Pro Gly Ala Val Gln Thr Leu Glu Gln Tyr Leu Phe Leu Tyr Arg 85 90 95

Ala Leu

<210> 324

<211> 355

<212> PRT

<213> Homo sapiens

| < 4.0 <i>(</i> |) | 2.4 | | | | | | | | | | | | | |
|----------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| |)> 32 | | | | | | | | | | | | | | |
| Met 1 | Ser | Arg | Gln | Leu 5 | Ser | Arg | Ala | Arg | Pro 10 | Ala | Thr | Val | Leu | Gly 15 | Ala |
| Met | Glu | Met | Gly 20 | Arg | Arg | Met | Asp | Ala 25 | Pro | Thr | Ser | Ala | Ala 30 | Val | Thr |
| Arg | Ala | Phe 35 | Leu | Glu | Arg | Gly | His 40 | Thr | Glu | Ile | Asp | Thr 45 | Ala | Phe | Val |
| Tyr | Ser 50 | Glu | Gly | Gln | Ser | Glu 55 | Thr | Ile | Leu | Gly | Gly 60 | Leu | Gly | Leu | Arg |
| Leu 65 | Gly | Gly | Ser | Asp | Cys 70 | Arg | Val | Lys | Ile | Asp 75 | Thr | Lys | Ala | Ile | Pro 80 |
| Leu | Phe | Gly | Asn | Ser 85 | Leu | Lys | Pro | Asp | Ser 90 | Leu | Arg | Phe | Gln | Leu 95 | Glu |
| Thr | Ser | Leu | Lys 100 | Arg | Leu | Gln | Cys | Pro 105 | Arg | Val | Asp | Leu | Phe 110 | Tyr | Leu |
| His | Met | Pro 115 | Asp | His | Ser | Thr | Pro 120 | Val | Glu | Glu | Thr | Leu 125 | Arg | Ala | Cys |
| His | Gln 130 | Leu | His | Gln | Glu | Gly 135 | Lys | Phe | Val | Glu | Leu 140 | Gly | Leu | Ser | Asn |
| Tyr 145 | Ala | Ala | Trp | Glu | Val 150 | Ala | Glu | Ile | Cys | Thr 155 | Leu | Cys | Lys | Ser | Asn 160 |
| Gly | Trp | Ile | Leu | Pro 165 | Thr | Val | Tyr | | Gly 170 | Met | Tyr | Asn | Ala | Ile 175 | Thr |
| Arg | Gln | Val | Glu 180 | Thr | Glu | Leu | Phe | Pro 185 | Cys | Leu | Arg | His | Phe 190 | Gly | Leu |
| Arg | Phe | Tyr 195 | Ala | Phe | Asn | Pro | Leu 200 | Ala | Asp | Gln | Ser | Pro 205 | Glu | Gly | Cys |
| Gly | Ser 210 | Phe | Trp | Gly | Thr | Leu 215 | Gly | Pro | Gly | Ala | Asp 220 | Cys | Cys | Phe | Pro |
| Ser 225 | Gly | Gly | Leu | Leu | Thr 230 | Gly | Lys | Tyr | Lys | Tyr 235 | Glu | Asp | Lys | Asn | Gly 240 |

Lys Gln Pro Val Gly Arg Phe Phe Gly Asn Thr Trp Ala Glu Met Tyr

| 245 | 250 | 255 |
|-----|-----|-----|
| 243 | 230 | 233 |

Arg Asn Arg Tyr Trp Lys Glu His His Phe Glu Gly Ile Ala Leu Val 260 265 270

Glu Lys Ala Leu Gln Ala Ala Tyr Gly Ala Ser Ala Pro Ser Met Thr 275 280 285

Ser Ala Thr Leu Arg Trp Met Tyr His His Ser Gln Leu Gln Gly Ala 290 295 300

His Gly Asp Ala Val Ile Leu Gly Met Ser Ser Leu Glu Gln Leu Glu 305 310 315 320

Gln Asn Leu Ala Ala Glu Glu Gly Pro Leu Glu Pro Ala Val Val 325 330 335

Asp Ala Phe Asn Gln Ala Trp His Leu Val Thr His Glu Cys Pro Asn 340 345 350

Tyr Phe Arg 355

<210> 325

<211> 331

<212> PRT

<213> Homo sapiens

<400> 325

Met Ser Arg Gln Leu Ser Arg Ala Arg Pro Ala Thr Val Leu Gly Ala 1 5 10 15

Met Glu Met Gly Arg Arg Met Asp Ala Pro Thr Ser Ala Ala Val Thr
20 25 30

Arg Ala Phe Leu Glu Arg Gly His Thr Glu Ile Asp Thr Ala Phe Val 35 40 45

Tyr Ser Asp Gly Gln Ser Glu Thr Ile Leu Gly Gly Leu Gly Leu Arg
50 55 60

Leu Gly Gly Ser Asp Cys Arg Val Lys Ile Asp Thr Lys Ala Ile Pro 65 70 75 80

Leu Phe Gly Asn Ser Leu Lys Pro Asp Ser Leu Arg Phe Gln Leu Glu 85 90 95

| Thr | Ser | Leu | Lys 100 | Arg | Leu | Gln | Cys | Pro 105 | Arg | Val | Asp | Leu | Phe 110 | Tyr | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Met | Pro 115 | Asp | His | Ser | Thr | Pro 120 | Val | Glu | Glu | Thr | Leu 125 | Arg | Ala | Cys |
| His | Gln 130 | Leu | His | Gln | Glu | Gly 135 | Lys | Phe | Met | Glu | Leu 140 | Gly | Leu | Ser | Asn |
| Tyr 145 | Ala | Ala | Trp | Glu | Val 150 | Ala | Glu | Ile | Cys | Thr 155 | Leu | Cys | Lys | Ser | Asn 160 |
| Gly | Trp | Ile | Leu | Pro 165 | Thr | Val | Tyr | Gln | Gly 170 | Met | Tyr | Asn | Ala | Ile 175 | Thr |
| Arg | Gln | Val | Glu 180 | Thr | Glu | Leu | Phe | Pro 185 | Cys | Leu | Arg | His | Phe 190 | Gly | Leu |
| Arg | Phe | Tyr 195 | Ala | Phe | Asn | Pro | Leu 200 | Ala | Gly | Gly | Leu | Leu 205 | Thr | Gly | Lys |
| Tyr | Lys 210 | Tyr | Glu | Asp | Lys | Asp 215 | Gly | Lys | Gln | Pro | Val 220 | Gly | Arg | Phe | Phe |
| Gly 225 | Asn | Thr | Trp | Ala | Glu 230 | Met | Tyr | Arg | Asn | Arg 235 | Tyr | Trp | Lys | Glu | His 240 |
| His | Phe | Glu | Gly | Ile 245 | Ala | Leu | Val | Glu | Lys 250 | Ala | Leu | Gln | Ala | Ala 255 | Tyr |
| Gly | Ala | Ser | Ala 260 | Pro | Ser | Met | Thr | Ser 265 | Ala | Thr | Leu | Arg | Trp 270 | Met | Tyr |
| His | His | Ser 275 | Gln | Leu | Gln | Gly | Ala 280 | His | Gly | Asp | Ala | Val 285 | Ile | Leu | Gly |
| Met | Ser 290 | Ser | Leu | Glu | Gln | Leu 295 | Glu | Gln | Asn | Leu | Ala 300 | Ala | Ala | Glu | Glu |
| Gly 305 | Pro | Leu | Glu | Pro | Ala 310 | Val | Val | Asp | Ala | Phe 315 | Asn | Gln | Ala | Trp | His 320 |
| Leu | Val | Ala | His | Glu 325 | Cys | Pro | Asn | Tyr | Phe 330 | Arg | | | | | |

<210> 326 <211> 331 <213> Homo sapiens

| <4 | n | <u>۵</u> > | 3 | 2 | 6 |
|----|---|------------|---|---|---|
| | | | | | |

- Met Ser Arg Gln Leu Ser Arg Ala Arg Pro Ala Thr Val Leu Gly Ala 1 5 10 15
- Met Glu Met Gly Arg Arg Met Asp Ala Pro Thr Ser Ala Ala Val Thr 20 25 30
- Arg Ala Phe Leu Glu Arg Gly His Thr Glu Ile Asp Thr Ala Phe Val $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$
- Tyr Ser Glu Gly Gln Ser Glu Thr Ile Leu Gly Gly Leu Gly Leu Arg
 50 55 60
- Leu Gly Gly Ser Asp Cys Arg Val Lys Ile Asp Thr Lys Ala Ile Pro 65 70 75 80
- Leu Phe Gly Asn Ser Leu Lys Pro Asp Ser Leu Arg Phe Gln Leu Glu 85 90 95
- Thr Ser Leu Lys Arg Leu Gln Cys Pro Arg Val Asp Leu Phe Tyr Leu 100 105 110
- His Met Pro Asp His Ser Thr Pro Val Glu Glu Thr Leu Arg Ala Cys
 115 120 125
- His Gln Leu His Gln Glu Gly Lys Phe Val Glu Leu Gly Leu Ser Asn 130 135 140
- Gly Trp Ile Leu Pro Thr Val Tyr Gln Gly Met Tyr Asn Ala Ile Thr 165 170 175
- Arg Gln Val Glu Thr Glu Leu Phe Pro Cys Leu Arg His Phe Gly Leu 180 185 190
- Arg Phe Tyr Ala Phe Asn Pro Leu Ala Gly Gly Leu Leu Thr Gly Lys
 195 200 205
- Tyr Lys Tyr Glu Asp Lys Asn Gly Lys Gln Pro Val Gly Arg Phe Phe 210 215 220
- Gly Asn Thr Trp Ala Glu Met Tyr Arg Asn Arg Tyr Trp Lys Glu His 225 230 235 240

His Phe Glu Gly Ile Ala Leu Val Glu Lys Ala Leu Gln Ala Ala Tyr Gly Ala Ser Ala Pro Ser Met Thr Ser Ala Thr Leu Arg Trp Met Tyr His His Ser Gln Leu Gln Gly Ala His Gly Asp Ala Val Ile Leu Gly Met Ser Ser Leu Glu Gln Leu Glu Gln Asn Leu Ala Ala Glu Glu Gly Pro Leu Glu Pro Ala Val Val Asp Ala Phe Asn Gln Ala Trp His Leu Val Thr His Glu Cys Pro Asn Tyr Phe Arg <210> 327 <211> 331 <212> PRT <213> Homo sapiens <400> 327 Met Ser Arg Gln Leu Ser Arg Ala Arg Pro Ala Thr Val Leu Gly Ala Met Glu Met Gly Arg Arg Met Asp Ala Pro Thr Ser Ala Ala Val Thr Arg Ala Phe Leu Glu Arg Gly His Thr Glu Ile Asp Thr Ala Phe Val Tyr Ser Glu Gly Gln Ser Glu Thr Ile Leu Gly Gly Leu Gly Leu Arg Leu Gly Gly Ser Asp Cys Arg Val Lys Ile Asp Thr Lys Ala Ile Pro

His Met Pro Asp His Ser Thr Pro Val Glu Glu Thr Leu Arg Ala Cys

Leu Phe Gly Asn Ser Leu Lys Pro Asp Ser Leu Arg Phe Gln Leu Glu

Thr Ser Leu Lys Arg Leu Gln Cys Pro Arg Val Asp Leu Phe Tyr Leu
100 105 110

His Gln Leu His Gln Glu Gly Lys Phe Val Glu Leu Gly Leu Ser Asn Tyr Ala Ala Trp Glu Val Ala Glu Ile Cys Thr Leu Cys Lys Ser Asn Gly Trp Ile Leu Pro Thr Val Tyr Gln Gly Met Tyr Asn Ala Ile Thr Arg Gln Val Glu Thr Glu Leu Phe Pro Cys Leu Arg His Phe Gly Leu Arg Phe Tyr Ala Phe Asn Pro Leu Ala Gly Gly Leu Leu Thr Gly Lys Tyr Lys Tyr Glu Asp Lys Asn Gly Lys Gln Pro Val Gly Arg Phe Phe Gly Asn Thr Trp Ala Glu Met Tyr Arg Asn Arg Tyr Trp Lys Glu His His Phe Glu Gly Ile Ala Leu Val Glu Lys Ala Leu Gln Ala Ala Tyr Gly Ala Ser Ala Pro Ser Met Thr Ser Ala Thr Leu Arg Trp Met Tyr His His Ser Gln Leu Gln Gly Ala His Gly Asp Ala Val Ile Leu Gly Met Ser Ser Leu Glu Gln Leu Glu Gln Asn Leu Ala Ala Glu Glu Gly Pro Leu Glu Pro Ala Val Val Asp Ala Phe Asn Gln Ala Trp His

Leu Val Thr His Glu Cys Pro Asn Tyr Phe Arg 325 330

<210> 328

<211> 330

<212> PRT

<213> Homo sapiens

<400> 328

| Met 1 | Ser | Arg | Pro | Pro 5 | Pro | Pro | Arg | Val | Ala 10 | Ser | Val | Leu | Gly | Thr 15 | Met |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Glu | Met | Gly | Arg 20 | Arg | Met | Asp | Ala | Pro 25 | Ala | Ser | Ala | Ala | Ala 30 | Val | Arg |
| Ala | Phe | Leu 35 | Glu | Arg | Gly | His | Thr 40 | Glu | Leu | Asp | Thr | Ala 45 | Phe | Met | Tyr |
| Ser | Asp 50 | Gly | Gln | Ser | Glu | Thr 55 | Ile | Leu | Gly | Gly | Leu 60 | Gly | Leu | Gly | Leu |
| Gly 65 | Gly | Gly | Asp | Cys | Arg 70 | Val | Lys | Ile | Ala | Thr 75 | Lys | Ala | Asn | Pro | Trp 80 |
| Asp | Gly | Lys | Ser | Leu 85 | Lys | Pro | Asp | Ser | Val 90 | Arg | Ser | Gln | Leu | Glu 95 | Thr |
| Ser | Leu | Lys | Arg 100 | Leu | Gln | Cys | Pro | Gln 105 | Val | Asp | Leu | Phe | Tyr 110 | Leu | His |
| Thr | Pro | Asp 115 | His | Gly | Thr | Pro | Val 120 | Glu | Glu | Thr | Leu | His 125 | Ala | Cys | Gln |
| Arg | Leu 130 | His | Gln | Glu | Gly | Lys 135 | Phe | Val | Glu | Leu | Gly 140 | Leu | Ser | Asn | Tyr |
| Ala 145 | Ser | Trp | Glu | Val | Ala 150 | Glu | Ile | Cys | Thr | Leu 155 | Cys | Lys | Ser | Asn | Gly 160 |
| Trp | Ile | Leu | Pro | Thr 165 | Val | Tyr | Gln | Gly | Met 170 | Tyr | Asn | Ala | Thr | Thr 175 | Arg |
| Gln | Val | Glu | Thr 180 | Glu | Leu | Phe | Pro | Cys 185 | Leu | Arg | His | Phe | Gly 190 | Leu | Arg |
| Phe | Tyr | Ala 195 | Tyr | Asn | Pro | Leu | Ala 200 | Gly | Gly | Leu | Leu | Thr 205 | Gly | Lys | Tyr |
| Lys | Tyr 210 | Glu | Asp | Lys | Asp | Gly 215 | Lys | Gln | Pro | Val | Gly 220 | Arg | Phe | Phe | Gly |
| Asn 225 | Ser | Trp | Ala | Glu | Thr 230 | Tyr | Arg | Asn | Arg | Phe 235 | Trp | Lys | Glu | His | His 240 |
| Phe | Glu | Ala | Ile | Ala 245 | Leu | Val | Glu | Lys | Ala 250 | Leu | Gln | Ala | Ala | Tyr 255 | Gly |

Ala Ser Ala Pro Ser Val Thr Ser Ala Ala Leu Arg Trp Met Tyr His His Ser Gln Leu Gln Gly Ala His Gly Asp Ala Val Ile Leu Gly Met Ser Ser Leu Glu Gln Leu Glu Gln Asn Leu Ala Ala Thr Glu Glu Gly Pro Leu Glu Pro Ala Val Val Asp Ala Phe Asn Gln Ala Trp His Leu Val Ala His Glu Cys Pro Asn Tyr Phe Arg <210> 329 <211> 306 <212> PRT <213> Homo sapiens <400> 329 Pro Ala Thr Val Leu Gly Ala Met Glu Met Gly Arg Arg Met Asp Ala Pro Thr Ser Ala Ala Val Thr Arg Ala Phe Leu Glu Arg Gly His Thr Glu Ile Asp Thr Ala Phe Leu Tyr Ser Asp Gly Gln Ser Glu Thr Ile Leu Gly Gly Leu Gly Leu Arg Met Gly Ser Ser Asp Cys Arg Val Lys Ile Ala Thr Lys Ala Asn Pro Trp Ile Gly Asn Ser Leu Lys Pro Asp Ser Val Arg Ser Gln Leu Glu Thr Ser Leu Lys Arg Leu Gln Cys Pro Arg Val Asp Leu Phe Tyr Leu His Ala Pro Asp His Ser Ala Pro Val Glu Glu Thr Leu Arg Ala Cys His Gln Leu His Gln Glu Gly Lys Phe

Val Glu Leu Gly Leu Ser Asn Tyr Ala Ala Trp Glu Val Ala Glu Ile

Cys Thr Leu Cys Lys Ser Asn Gly Trp Ile Leu Pro Thr Val Tyr Gln 145 150 155 Gly Met Tyr Ser Ala Thr Thr Arg Gln Val Glu Thr Glu Leu Phe Pro 165 170 175 Cys Leu Arg His Phe Gly Leu Arg Phe Tyr Ala Tyr Asn Pro Leu Ala 180 185 Asp Gln Ser Pro Glu Gly Cys Gly Ser Phe Trp Gly Thr Leu Gly Pro 195 200 205 Gly Ala Asp Cys Cys Leu Pro Ala Gly Gly Leu Leu Thr Gly Lys Tyr 215 220 Lys Tyr Glu Asp Lys Asp Gly Lys Gln Pro Val Gly Arg Phe Phe Gly 230 235 Thr Gln Trp Ala Glu Ile Tyr Arg Asn Gln Phe Trp Lys Glu His His 245 250 Phe Glu Gly Ile Ala Leu Val Glu Lys Ala Leu Gln Ala Ala Tyr Gly 260 265 Ala Ser Ala Pro Ser Met Thr Ser Ala Ala Leu Arg Trp Met Tyr His 275 280 285 His Ser Gln Leu Gln Gly Ala His Gly Asp Ala Val Ile Leu Gly Met 290 295 300 Ser Ser 305 <210> 330 <211> 245 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Aldo/keto reductase family domain sequence <400> 330

Pro Leu Leu Gly Leu Gly Thr Trp Lys Thr Pro Gly Arg Val Asp Asp

15

5

Glu Glu Ala Phe Glu Ala Val Lys Ala Ala Leu Asp Ala Gly Tyr Arg
20 25 30

His Phe Asp Thr Ala Glu Ile Tyr Gly Asn Glu Glu Glu Val Gly Glu 35 40 45

Ala Ile Lys Glu Ala Leu Phe Glu Gly Ser Gly Val Val Arg Glu Asp 50 55 60

Ile Phe Ile Thr Ser Lys Leu Trp Asn Thr Phe His Ser Pro Lys His 65 70 75 80

Val Arg Glu Ala Leu Glu Lys Ser Leu Lys Arg Leu Gly Leu Asp Tyr 85 90 95

Val Asp Leu Tyr Leu Ile His Trp Pro Asp Pro Leu Lys Pro Gly Asp
100 105 110

Asp Val Pro Ile Glu Glu Thr Trp Lys Ala Leu Glu Lys Leu Val Asp 115 120 125

Glu Gly Lys Val Arg Ser Ile Gly Val Ser Asn Phe Ser Ala Glu Gln 130 135 140

Gln Val Glu Tyr His Pro Tyr Leu Arg Gln Asp Glu Leu Arg Lys Phe 165 170 175

Cys Lys Lys His Gly Ile Gly Val Thr Ala Tyr Ser Pro Leu Gly Ser 180 185 190

Gly Leu Leu Asp Lys Phe Trp Ser Glu Leu Gly Ser Pro Glu Leu Leu 195 200 205

Glu Asp Pro Ala Leu Lys Lys Ile Ala Glu Lys Tyr Gly Lys Thr Pro 210 215 220

Ala Gln Val Ala Leu Arg Trp Val Leu Gln Arg Gly Val Ser Val Ile 225 230 235 240

Pro Lys Ser Ser Thr 245

<210> 331 <211> 768

<213> Mus musculus

| < 4 | 0 | 0> | 3 | 3 | 1 |
|-----|---|----|---|---|---|
| | | | | | |

- Met Lys Leu Leu Trp Gln Ala Lys Met Ser Ser Ile Gln Asp Trp Gly
 1 5 10 15
- Glu Glu Val Glu Glu Gly Ala Val Tyr His Val Thr Leu Lys Arg Val
 20 25 30
- Gln Ile Gln Gln Ala Ala Asn Lys Gly Ala Arg Trp Leu Gly Val Glu 35 40 45
- Gly Asp Gln Leu Pro Pro Gly His Thr Val Ser Gln Tyr Glu Thr Cys
 50 55 60
- Lys Ile Arg Thr Ile Lys Ala Gly Thr Leu Glu Lys Leu Val Glu Asn 65 70 75 80
- Leu Leu Thr Ala Phe Gly Asp Asn Asp Phe Thr Tyr Ile Ser Ile Phe 85 90 95
- Leu Ser Thr Tyr Arg Gly Phe Ala Ser Thr Lys Glu Val Leu Glu Leu
 100 105 110
- Leu Leu Asp Arg Tyr Gly Asn Leu Thr Ser Pro Asn Cys Glu Glu Asp 115 120 125
- Gly Ser Gln Ser Ser Glu Ser Lys Met Val Ile Arg Asn Ala Ile 130 135 140
- Ala Ser Ile Leu Arg Ala Trp Leu Asp Gln Cys Ala Glu Asp Phe Arg 145 150 155 160
- Glu Pro Pro His Phe Pro Cys Leu Gln Lys Leu Leu Asp Tyr Leu Thr 165 170 175
- Arg Met Met Pro Gly Ser Asp Pro Glu Arg Arg Ala Gln Asn Leu Leu 180 185 190
- Glu Gln Phe Gln Lys Gln Glu Val Glu Thr Asp Asn Gly Leu Pro Asn 195 200 205
- Thr Ile Ser Phe Ser Leu Glu Glu Glu Glu Glu Leu Glu Gly Gly Glu 210 215 220
- Ser Ala Glu Phe Thr Cys Phe Ser Glu Asp Leu Val Ala Glu Gln Leu 225 230 235 240

Thr Tyr Met Asp Ala Gln Leu Phe Lys Lys Val Val Pro His His Cys Leu Gly Cys Ile Trp Ser Arg Arg Asp Lys Lys Glu Asn Lys His Leu Ala Pro Thr Ile Arg Ala Thr Ile Ser Gln Phe Asn Thr Leu Thr Lys Cys Val Val Ser Thr Ile Leu Gly Gly Lys Glu Leu Lys Thr Gln Gln Arg Ala Lys Ile Ile Glu Lys Trp Ile Asn Ile Ala His Glu Cys Arg Leu Leu Lys Asn Phe Ser Ser Leu Arg Ala Ile Val Ser Ala Leu Gln Ser Asn Ser Ile Tyr Arg Leu Lys Lys Thr Trp Ala Ala Val Pro Arg Asp Arg Met Leu Met Phe Glu Glu Leu Ser Asp Ile Phe Ser Asp His Asn Asn His Leu Thr Ser Arg Glu Leu Leu Met Lys Glu Gly Thr Ser Lys Phe Ala Asn Leu Asp Ser Ser Val Lys Glu Asn Gln Lys Arg Thr Gln Arg Arg Leu Gln Leu Gln Lys Asp Met Gly Val Met Gln Gly Thr Val Pro Tyr Leu Gly Thr Phe Leu Thr Asp Leu Thr Met Leu Asp Thr Ala Leu Gln Asp Tyr Ile Glu Gly Gly Leu Ile Asn Phe Glu Lys Arg Arg Arg Glu Phe Glu Val Ile Ala Gln Ile Lys Leu Gln Ser Ala Cys Asn Ser Tyr Cys Met Thr Pro Asp Gln Lys Phe Ile Gln Trp Phe Gln Arg Gln Gln Leu Leu Thr Glu Glu Ser Tyr Ala Leu Ser Cys

Glu Ile Glu Ala Ala Asp Ala Ser Thr Thr Ser Pro Lys Pro Arg Lys Ser Met Val Lys Arg Leu Ser Leu Leu Phe Leu Gly Ser Asp Met Ile Thr Ser Pro Thr Pro Thr Lys Glu Gln Pro Lys Ser Thr Ala Ser Gly Ser Ser Gly Glu Ser Met Asp Ser Val Ser Val Ser Ser Cys Glu Ser Asn His Ser Glu Ala Glu Glu Gly Ser Ile Thr Pro Met Asp Thr Pro Asp Glu Pro Gln Lys Lys Leu Ser Glu Ser Ser Ser Ser Cys Ser Ser Ile His Ser Met Asp Thr Asn Ser Ser Gly Met Ser Ser Leu Ile Asn Pro Leu Ser Ser Pro Pro Ser Cys Asn Asn Pro Lys Ile His Lys Arg Ser Val Ser Val Thr Ser Ile Thr Ser Thr Val Leu Pro Pro Val Tyr Asn Gln Gln Asn Glu Asp Thr Cys Ile Ile Arg Ile Ser Val Glu Asp Asn Asn Gly Asn Met Tyr Lys Ser Ile Met Leu Thr Ser Gln Asp Lys Thr Pro Ala Val Ile Gln Arg Ala Met Leu Lys His Asn Leu Asp Ser Asp Pro Ala Glu Glu Tyr Glu Leu Val Gln Val Ile Ser Glu Asp Lys Glu Leu Val Ile Pro Asp Ser Ala Asn Val Phe Tyr Ala Met Asn Ser Gln Val Asn Phe Asp Phe Ile Leu Arg Lys Lys Asn Ser Met Glu Glu Gln Val Lys Leu Arg Ser Arg Thr Ser Leu Thr Leu Pro Arg

Thr Ala Lys Arg Gly Cys Trp Ser Xaa Arg His Ser Lys Ile Thr Leu 755 760 765

<210> 332

<211> 709

<212> PRT

<213> Mus musculus

<400> 332

Met Glu Arg Thr Ala Gly Lys Glu Leu Ala Leu Ala Pro Leu Gln Asp
1 5 10 15

Trp Gly Glu Glu Thr Glu Asp Gly Ala Val Tyr Ser Val Ser Leu Arg
20 25 30

Arg Gln Arg Ser Gln Arg Ser Thr Pro Glu Arg Ser Gly Glu Gly Gln
35 40 45

Thr Pro Ile Pro Ala Thr Asp Thr Phe Leu His Tyr Arg Thr Ser Lys 50 55 60

Val Arg Ala Leu Arg Ala Ala Arg Leu Glu Arg Leu Val His Glu Leu 65 70 75 80

Val Ser Gly Asp Arg Glu Gln Asp Pro Gly Phe Val Pro Ala Phe Leu 85 90 95

Ala Thr His Arg Ala Phe Val Pro Thr Ala Arg Val Leu Gly Phe Leu
100 105 110

Lys Arg Thr Glu Gly Gln Asp Leu Asn Phe Ser Lys Asn Leu Arg Ala 130 135 140

Val Val Ser Val Leu Gly Ser Trp Leu Arg Asn His Pro Gln Asp Phe 145 150 155 160

Arg Asp Pro Pro Asp His Gln Asn Leu Gly Asn Val Arg Ile Phe Leu 165 170 175

Gly Trp Val Ala Pro Gly Gly Ala Glu Ala Arg Glu Ala Glu Lys Leu

| | | | 180 | | | | | 185 | | | | | 190 | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Glu | Asp 195 | Phe | Leu | Lys | Glu | Ala 200 | Lys | Gly | Glu | Gln | Thr 205 | Glu | Glu | Glu |
| Lys | Arg 210 | Leu | Ala | Trp | Ser | Gly 215 | Pro | Pro | Arg | Ile | Ala 220 | Gln | Thr | Pro | Gly |
| Ser 225 | Glu | Phe | Ala | Glu | Asp 230 | Суѕ | Val | Glu | Glu | Glu 235 | Gly | Pro | Ser | Ser | Glu 240 |
| Gly | Pro | Glu | Leu | Leu 245 | Asp | Phe | Ser | Val | Asp 250 | Asp | Val | Ala | Glu | Gln 255 | Leu |
| Thr | Leu | Met | Asp 260 | Val | Glu | Leu | Phe | Leu 265 | Arg | Val | Arg | Ser | Cys 270 | Glu | Cys |
| Leu | Gly | Ser 275 | Met | Trp | Ser | Gln | Arg 280 | Asp | Arg | Pro | Gly | Ala 285 | Ala | Gly | Ile |
| Ser | Pro 290 | Thr | Val | Arg | Ala | Thr 295 | Val | Ala | Gln | Phe | Asn 300 | Thr | Val | Thr | Gly |
| Cys 305 | Val | Leu | Gly | Ser | Val 310 | Leu | Ala | Ala | Pro | Gly 315 | Leu | Ala | Ala | Ser | Gln 320 |
| Arg | Ala | Gln | Arg | Ile 325 | Glu | Lys | Trp | Ile | Arg 330 | Ile | Ala | Gln | Arg | Cys 335 | Arg |
| Glu | Leu | Arg | Asn 340 | Phe | Ser | Ser | Leu | Arg 345 | Ala | Ile | Leu | Ser | Ala 350 | Leu | Gln |
| Ser | Asn | Pro 355 | Ile | Tyr | Arg | Leu | Lys 360 | Arg | Ser | Trp | Gly | Ala 365 | Val | Ser | Arg |
| Glu | Pro 370 | Leu | Ser | Val | Phe | Arg 375 | Lys | Leu | Ser | Gln | Ile 380 | Phe | Ser | Asp | Glu |
| Asp 385 | Asn | His | Leu | Ser | Ser 390 | Arg | Ala | Ile | Leu | Ser 395 | Gln | Glu | Glu | Thr | Thr 400 |
| Glu | Asp | Asp | Asp | Cys 405 | Pro | Ser | Gly | Ser | Leu 410 | Pro | Ser | Lys | Leu | Pro 415 | Pro |
| Gly | Pro | Val | Pro 420 | Tyr | Leu | Gly | Thr | Phe 425 | Leu | Thr | Asp | Leu | Val 430 | Met | Leu |

Asp Thr Ala Leu Pro Asp Thr Leu Lys Gly Asn Leu Ile Asn Phe Glu

| 445 |
|-----|
| |

| Lys | Arg 450 | Arg | Lys | Glu | Trp | Glu 455 | Ile | Leu | Ala | Arg | Ile 460 | Gln | Gln | Leu | Gln |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gln 465 | Arg | Cys | Gln | Arg | Tyr 470 | Ser | Leu | Ser | Pro | Arg 475 | Pro | Pro | Ile | Leu | Ala 480 |
| Ala | Leu | Arg | Ala | Gln 485 | Arg | Gln | Leu | Ser | Glu 490 | Glu | Gln | Ser | Tyr | Arg 495 | Val |
| Ser | Arg | Val | Ile 500 | Glu | Pro | Pro | Ala | Ala 505 | Ser | Cys | Pro | Ser | Ser 510 | Pro | Arg |
| Ile | Arg | Arg 515 | Arg | Ile | Ser | Leu | Thr 520 | Lys | Arg | Leu | Ser | Ala 525 | Lys | Leu | Ser |
| Arg | Glu 530 | Lys | Asn | Ser | Ser | Pro 535 | Gly | Gly | Ser | Pro | Gly 540 | Asp | Pro | Ser | Ser |
| Pro 545 | Thr | Ser | Ser | Val | Ser 550 | Pro | Gly | Ser | Pro | Pro 555 | Ser | Ser | Pro | Arg | Asn 560 |
| Arg | Glu | Pro | Pro | Pro 565 | Pro | Gly | Ser | Pro | Pro 570 | Ala | Ser | Pro | Gly | Pro 575 | Gln |
| Ser | Pro | Ser | Thr 580 | Lys | Leu | Ser | Leu | Thr 585 | Met | Asp | Pro | Pro | Gly 590 | Pro | Trp |
| Pro | Val | Thr 595 | Leu | Thr | Pro | Ser | Ser 600 | Ser | Arg | Val | Pro | Leu 605 | Leu | Gly | Gln |
| Gln | Thr 610 | Ser | Glu | Ala | Arg | Val 615 | Ile | Arg | Val | Ser | Ile 620 | Asn | Asn | Asn | His |
| Gly 625 | Asn | Leu | Tyr | Arg | Ser 630 | Ile | Leu | Leu | Thr | Cys 635 | Gln | Asp | Lys | Ala | Pro 640 |
| Ser | Val | Val | Gln | Arg 645 | Ala | Leu | Glu | Lys | His 650 | Asn | Val | Pro | Gln | Pro 655 | Trp |
| Ala | Arg | Asp | Tyr 660 | Gln | Leu | Phe | Gln | Val 665 | Leu | Pro | Gly | Asp | Arg 670 | Glu | Leu |
| Leu | Ile | Pro 675 | Asp | Gly | Ala | Asn | Val 680 | Phe | Tyr | Ala | Met | Ser 685 | Pro | Ala | Ala |
| Pro | Glv | Asp | Phe | Leu | Leu | Ara | Ara | Lvs | Glu | Glv | Thr | Glv | His | Thr | Leu |

690 695 700

Ser Ala Ser Pro Thr 705

<210> 333

<211> 343

<212> PRT

<213> Mus musculus

<400> 333

Met Ala Pro Cys Thr Ala Ser Pro Cys Gly Gly Ser Ala Ala Ser Ala 1 5 10 15

Arg Pro Gln Arg Gly Leu Glu Lys Ala Arg Val Asp Ser Lys Arg Thr
20 25 30

Glu Gly Gln Asp Leu Asn Phe Ser Lys Asn Leu Arg Ala Val Val Ser 35 40 45

Val Leu Gly Ser Trp Leu Arg Asn His Pro Gln Asp Phe Arg Asp Pro 50 55 60

Pro Asp His Gln Asn Leu Gly Asn Val Arg Ile Phe Leu Gly Trp Ala 65 70 75 80

Ala Pro Gly Gly Ala Glu Ala Arg Glu Ala Glu Lys Leu Leu Glu Asp 85 90 95

Phe Leu Lys Glu Ala Lys Gly Glu Gln Thr Glu Glu Glu Lys Arg Leu
100 105 110

Ala Trp Ser Gly Pro Pro Arg Ile Ala Gln Thr Pro Gly Ser Glu Phe
115 120 125

Ala Glu Asp Cys Val Glu Glu Glu Gly Pro Ser Ser Glu Gly Pro Glu 130 135 140

Asp Val Glu Leu Phe Leu Arg Val Arg Ser Cys Glu Cys Leu Gly Ser 165 170 175

Met Trp Ser Gln Arg Asp Arg Pro Gly Ala Ala Gly Ile Ser Pro Thr 180 185 190

Gly Ser Val Leu Ala Ala Pro Gly Leu Ala Ala Ser Gln Lys Ala Gln Arg Ile Glu Lys Trp Ile Arg Ile Ala Gln Arg Cys Arg Glu Leu Arg Asn Phe Ser Ser Leu Arg Ala Ile Leu Ser Ala Leu Gln Ser Asn Pro Ile Tyr Arg Leu Lys Arg Ser Trp Gly Ala Val Ser Arg Glu Pro Leu Ser Val Phe Arg Lys Leu Ser Gln Ile Phe Ser Asp Glu Asp Asn His Leu Ser Ser Arg Ala Ile Leu Ser Gln Glu Glu Thr Thr Glu Asp Asp Asp Cys Pro Ser Gly Ser Leu Pro Ser Lys Leu Pro Pro Gly Pro Val Pro Tyr Leu Gly Thr Phe Leu Thr Asp Leu Val Met Leu Asp Thr Ala Leu Pro Asp Thr Leu Lys Val <210> 334 <211> 343 <212> PRT <213> Homo sapiens <400> 334 Met Ala Pro Cys Thr Ala Ser Pro Cys Gly Gly Ser Ala Ala Ser Ala Arg Pro Gln Arg Gly Leu Glu Lys Ala Arg Val Asp Ser Lys Arg Thr Glu Gly Gln Asp Leu Asn Phe Ser Lys Asn Leu Arg Ala Val Val Ser Val Leu Gly Ser Trp Leu Arg Asn His Pro Gln Asp Phe Arg Asp Pro

Val Arg Ala Thr Val Ala Gln Phe Asn Thr Val Thr Gly Cys Val Leu

| Pro 65 | Asp | His | Gln | Asn | Leu 70 | Gly | Asn | Val | Arg | Ile 75 | Phe | Leu | Gly | Trp | Ala 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Pro | Gly | Gly | Ala 85 | Glu | Ala | Arg | Glu | Ala 90 | Glu | Lys | Leu | Leu | Glu 95 | Asp |
| Phe | Leu | Lys | Glu 100 | Ala | Lys | Gly | Glu | Gln 105 | Thr | Glu | Glu | Glu | Lys 110 | Arg | Leu |
| Ala | Trp | Ser 115 | Gly | Pro | Pro | Arg | Ile 120 | Ala | Gln | Thr | Pro | Gly 125 | Ser | Glu | Phe |
| Ala | Glu 130 | Asp | Cys | Val | Glu | Glu 135 | Glu | Gly | Pro | Ser | Ser 140 | Glu | Gly | Pro | Glu |
| Leu 145 | Leu | Asp | Phe | Ser | Val 150 | Asp | Asp | Val | Ala | Glu 155 | Gln | Leu | Thr | Leu | Met 160 |
| Asp | Val | Glu | Leu | Phe 165 | Leu | Arg | Val | Arg | Ser 170 | Cys | Glu | Cys | Leu | Gly 175 | Ser |
| Met | Trp | Ser | Gln 180 | Arg | Asp | Arg | Pro | Gly 185 | Ala | Ala | Gly | Ile | Ser 190 | Pro | Thr |
| Val | Arg | Ala 195 | Thr | Val | Ala | Gln | Phe 200 | Asn | Thr | Val | Thr | Gly 205 | Cys | Val | Leu |
| Gly | Ser 210 | Val | Leu | Ala | Ala | Pro 215 | Gly | Leu | Ala | Ala | Ser 220 | Gln | Lys | Ala | Gln |
| Arg 225 | Ile | Glu | Lys | Trp | Ile 230 | Arg | Ile | Ala | Gln | Arg 235 | Cys | Arg | Glu | Leu | Arg 240 |
| Asn | Phe | Ser | Ser | Leu 245 | Arg | Ala | Ile | Leu | Ser 250 | Ala | Leu | Gln | Ser | Asn 255 | Pro |
| Ile | Tyr | Arg | Leu 260 | Lys | Arg | Ser | Trp | Gly 265 | Ala | Val | Ser | Arg | Glu 270 | Pro | Leu |
| Ser | Val | Phe 275 | Arg | Lys | Leu | Ser | Gln 280 | Ile | Phe | Ser | Asp | Glu 285 | Asp | Asn | His |
| Leu | Ser 290 | Ser | Arg | Ala | Ile | Leu 295 | Ser | Gln | Glu | Glu | Thr 300 | Thr | Glu | Asp | Asp |
| Asp 305 | Cys | Pro | Ser | Gly | Ser 310 | Leu | Pro | Ser | Lys | Leu 315 | Pro | Pro | Gly | Pro | Val 320 |

Pro Tyr Leu Gly Thr Phe Leu Thr Asp Leu Val Met Leu Asp Thr Ala 325 330 335

Leu Pro Asp Thr Leu Lys Val

<210> 335

<211> 709

<212> PRT

<213> Mus musculus

<400> 335

Met Glu Arg Thr Ala Gly Lys Glu Leu Ala Leu Ala Pro Leu Gln Asp 1 5 10 15

Trp Gly Glu Glu Thr Glu Asp Gly Ala Val Tyr Ser Val Ser Leu Arg
20 25 30

Arg Gln Arg Ser Gln Arg Ser Thr Pro Glu Arg Ser Gly Glu Gly Gln
35 40 45

Thr Pro Ile Pro Ala Thr Asp Thr Phe Leu His Tyr Arg Thr Ser Lys 50 55 60

Val Arg Ala Leu Arg Ala Ala Arg Leu Glu Arg Leu Val His Glu Leu 65 70 75 80

Val Ser Gly Asp Arg Glu Gln Asp Pro Gly Phe Val Pro Ala Phe Leu 85 90 95

Ala Thr His Arg Ala Phe Val Pro Thr Ala Arg Val Leu Gly Phe Leu
100 105 110

Leu Pro Pro Pro Pro Pro Pro Pro Pro Pro Ala Gly Val Asp Ser 115 120 125

Lys Arg Thr Glu Gly Gln Asp Leu Asn Phe Ser Lys Asn Leu Arg Ala 130 135 140

Val Val Ser Val Leu Gly Ser Trp Leu Arg Asn His Pro Gln Asp Phe 145 150 155 160

Arg Asp Pro Pro Asp His Gln Asn Leu Gly Asn Val Arg Ile Phe Leu 165 170 175

Gly Trp Ala Ala Pro Gly Gly Ala Glu Ala Arg Glu Ala Glu Lys Leu

| | | | 180 | | | | | 185 | | | | | 190 | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Glu | Asp 195 | Phe | Leu | Lys | Glu | Ala 200 | Lys | Gly | Glu | Gln | Thr 205 | Glu | Glu | Glu |
| Lys | Arg 210 | Leu | Ala | Trp | Ser | Gly 215 | Pro | Pro | Arg | Ile | Ala 220 | Gln | Thr | Pro | Gly |
| Ser 225 | Glu | Phe | Ala | Glu | Asp 230 | Cys | Val | Glu | Glu | Glu 235 | Gly | Pro | Ser | Ser | Glu 240 |
| Gly | Pro | Glu | Leu | Leu 245 | Asp | Phe | Ser | Val | Asp 250 | Asp | Val | Ala | Glu | Gln 255 | Leu |
| Thr | Leu | Met | Asp 260 | Val | Glu | Leu | Phe | Leu 265 | Arg | Val | Arg | Ser | Cys 270 | Glu | Cys |
| Leu | Gly | Ser 275 | Met | Trp | Ser | Gln | Arg 280 | Asp | Arg | Pro | Gly | Ala 285 | Ala | Gly | Ile |
| Ser | Pro 290 | Thr | Val | Arg | Ala | Thr 295 | Val | Ala | Gln | Phe | Asn 300 | Thr | Val | Thr | Gly |
| Cys 305 | Val | Leu | Gly | Ser | Val 310 | Leu | Ala | Ala | Pro | Gly 315 | Leu | Ala | Ala | Ser | Gln 320 |
| Arg | Ala | Gln | Arg | Ile 325 | Glu | Lys | Trp | Ile | Arg 330 | Ile | Ala | Gln | Arg | Cys 335 | Arg |
| Glu | Leu | Arg | Asn 340 | Phe | Ser | Ser | Leu | Arg 345 | Ala | Ile | Leu | Ser | Ala 350 | Leu | Gln |
| Ser | Asn | Pro 355 | Ile | Tyr | Arg | Leu | Lys 360 | Arg | Ser | Trp | Gly | Ala 365 | Val | Ser | Arg |
| Glu | Pro 370 | Leu | Ser | Val | Phe | Arg 375 | Lys | Leu | Ser | Gln | Ile 380 | Phe | Ser | Asp | Glu |
| Asp 385 | Asn | His | Leu | Ser | Ser 390 | Arg | Ala | Ile | Leu | Ser 395 | Gln | Glu | Glu | Thr | Thr 400 |
| Glu | Asp | Asp | Asp | Cys 405 | Pro | Ser | Gly | Ser | Leu 410 | Pro | Ser | Lys | Leu | Pro 415 | Pro |
| Gly | Pro | Val | Pro 420 | Tyr | Leu | Gly | Thr | Phe 425 | Leu | Thr | Asp | Leu | Val 430 | Met | Leu |

Asp Thr Ala Leu Pro Asp Thr Leu Lys Gly Asn Leu Ile Asn Phe Glu

| 435 | 440 | 445 |
|-----|-----|-----|
| | | |

| Lys | Arg 450 | Arg | Lys | Glu | Trp | Glu 455 | Ile | Leu | Ala | Arg | Ile 460 | Gln | Gln | Leu | Gln |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gln 465 | Arg | Cys | Gln | Arg | Tyr 470 | Ser | Leu | Ser | Pro | Arg 475 | Pro | Pro | Ile | Leu | Ala 480 |
| Ala | Leu | Arg | Ala | Gln 485 | Arg | Gln | Leu | Ser | Glu 490 | Glu | Gln | Ser | Tyr | Arg 495 | Val |
| Ser | Arg | Val | Ile 500 | Glu | Pro | Pro | Ala | Ala 505 | Ser | Cys | Pro | Ser | Ser 510 | Pro | Arg |
| Ile | Arg | Arg 515 | Arg | Ile | Ser | Leu | Thr 520 | Lys | Arg | Leu | Ser | Ala 525 | Lys | Leu | Ser |
| Arg | Glu 530 | Lys | Asn | Ser | Ser | Pro 535 | Gly | Gly | Ser | Pro | Gly 540 | Asp | Pro | Ser | Ser |
| Pro 545 | Thr | Ser | Ser | Val | Ser 550 | Pro | Gly | Ser | Pro | Pro 555 | Ser | Ser | Pro | Arg | Asn 560 |
| Arg | Glu | Pro | Pro | Pro 565 | Pro | Gly | Ser | Pro | Pro 570 | Ala | Ser | Pro | Gly | Pro 575 | Gln |
| Ser | Pro | Ser | Thr 580 | Lys | Leu | Ser | Leu | Thr 585 | Met | Asp | Pro | Pro | Gly 590 | Pro | Trp |
| Pro | Val | Thr 595 | Leu | Thr | Pro | Ser | Ser 600 | Ser | Arg | Val | Pro | Leu 605 | Leu | Gly | Gln |
| Gln | Thr 610 | Ser | Glu | Ala | Arg | Val 615 | Ile | Arg | Val | Ser | Ile 620 | Asn | Asn | Asn | His |
| Gly 625 | Asn | Leu | Tyr | Arg | Ser 630 | Ile | Leu | Leu | Thr | Cys 635 | Gln | Asp | Lys | Ala | Pro 640 |
| Ser | Val | Val | Gln | Arg 645 | Ala | Leu | Glu | Lys | His 650 | Asn | Val | Pro | Gln | Pro 655 | Trp |
| Ala | Arg | Asp | Tyr 660 | Gln | Leu | Phe | Gln | Val 665 | Leu | Pro | Gly | Asp | Arg 670 | Glu | Leu |
| Leu | Ile | Pro 675 | Asp | Gly | Ala | Asn | Val 680 | Phe | Tyr | Ala | Met | Ser 685 | Pro | Ala | Ala |
| Pro | Gly | Asp | Phe | Leu | Leu | Arg | Arg | Lys | Glu | Gly | Thr | Gly | His | Thr | Leu |

690 695 700

Ser Ala Ser Pro Thr 705

<210> 336

<211> 261

<212> PRT

<213> Homo sapiens

<400> 336

Leu Leu Asp Phe Ser Val Asp Glu Val Ala Glu Gln Leu Thr Leu Ile 1 5 10 15

Asp Leu Glu Leu Phe Ser Lys Val Arg Leu Tyr Glu Cys Leu Gly Ser 20 25 30

Val Trp Ser Gln Arg Asp Arg Pro Gly Ala Ala Gly Ala Ser Pro Thr 35 40 45

Val Arg Ala Thr Val Ala Gln Phe Asn Thr Val Thr Gly Cys Val Leu 50 55 60

Gly Ser Val Leu Gly Ala Pro Gly Leu Ala Ala Pro Gln Arg Ala Gln 65 70 75 80

Arg Leu Glu Lys Trp Ile Arg Ile Ala Gln Arg Cys Arg Glu Leu Arg 85 90 95

Asn Phe Ser Ser Leu Arg Ala Ile Leu Ser Ala Leu Gln Ser Asn Pro 100 105 110

Ile Tyr Arg Leu Lys Arg Ser Trp Gly Ala Val Ser Arg Glu Pro Leu 115 120 125

Ser Thr Phe Arg Lys Leu Ser Gln Ile Phe Ser Asp Glu Asn Asn His 130 135 140

Leu Ser Ser Arg Glu Ile Leu Phe Gln Glu Glu Ala Thr Glu Gly Ser 145 150 155 160

Gln Glu Glu Asp Asn Thr Pro Gly Ser Leu Pro Ser Lys Pro Pro Pro 165 170 175

Gly Pro Val Pro Tyr Leu Gly Thr Phe Leu Thr Asp Leu Val Met Leu 180 185 190 Asp Thr Ala Leu Pro Asp Met Leu Glu Gly Asp Leu Ile Asn Phe Glu
195 200 205

Lys Arg Arg Lys Glu Trp Glu Ile Leu Ala Arg Ile Gln Gln Leu Gln 210 215 220

Arg Arg Cys Gln Ser Tyr Thr Leu Ser Pro His Pro Pro Ile Leu Ala 225 230 235 240

Ala Leu His Ala Gln Asn Gln Leu Thr Glu Glu Gln Ser Tyr Arg Leu 245 250 255

Ser Arg Val Ile Glu 260

<210> 337

<211> 239

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: RasGEF domain sequence

<400> 337

Leu Leu Leu Leu Asp Pro Lys Glu Leu Ala Glu Gln Leu Thr Leu Leu 1 5 10 15

Asp Phe Glu Leu Phe Arg Lys Ile Asp Pro Ser Glu Leu Leu Gly Ser 20 . 25 30

Val Trp Gly Lys Arg Ser Lys Lys Ser Pro Ser Pro Leu Asn Leu Glu 35 40 45

Arg Phe Ile Glu Arg Phe Asn Glu Val Ser Asn Trp Val Ala Thr Glu 50 55 60

Ile Leu Lys Gln Thr Thr Pro Lys Asp Arg Ala Glu Leu Leu Ser Lys 65 70 75 80

Phe Ile Gln Val Ala Lys His Cys Arg Glu Leu Asn Asn Phe Asn Ser 85 90 95

Leu Met Ala Ile Val Ser Ala Leu Ser Ser Ser Pro Ile Ser Arg Leu 100 105 110

Lys Lys Thr Trp Glu Lys Leu Pro Ser Lys Tyr Lys Leu Phe Glu

| | | 115 | | | | | 120 | | | | | 125 | | | |
|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|------------|-----|-----|-----|-----|
| Glu | Leu 130 | Glu | Glu | Leu | Leu | Asp 135 | Pro | Ser | Arg | Asn | Phe 140 | Lys | Asn | Tyr | Arg |
| Glu | Ala | Leu | Ser | Ser | Cys | Asn | Leu | Pro | Pro | Cys | Ile | Pro | Phe | Leu | Gly |

145 150 155 160

Val Leu Leu Lys Asp Leu Thr Phe Ile Asp Glu Gly Asn Pro Asp Phe 165 170 175

Leu Lys Asn Gly Leu Val Asn Phe Glu Lys Arg Arg Lys Ile Ala Lys 180 185 190

Ile Leu Arg Glu Ile Arg Gln Leu Gln Ser Gln Pro Tyr Asn Leu Arg 195 200 205

Pro Asn Arg Ser Asp Ile Gln Ser Leu Leu Gln Gln Ser Leu Asp Ser 210 215 220

Leu Pro Glu Glu Asn Glu Leu Tyr Glu Leu Ser Leu Arg Ile Glu 225 230 235

<210> 338

<211> 211

<212> PRT

<213> Homo sapiens

<400> 338

Leu Asp Phe Ser Val Asp Glu Val Ala Glu Gln Leu Thr Leu Ile Asp 1 5 10 15

Leu Glu Leu Phe Ser Lys Val Arg Leu Tyr Glu Cys Leu Gly Ser Val $20 \hspace{1.5cm} 25 \hspace{1.5cm} 30$

Trp Ser Gln Arg Asp Arg Pro Gly Ala Ala Gly Ala Ser Pro Thr Val 35 40 45

Arg Ala Thr Val Ala Gln Phe Asn Thr Val Thr Gly Cys Val Leu Gly 50 55 60

Ser Val Leu Gly Ala Pro Gly Leu Ala Ala Pro Gln Arg Ala Gln Arg 65 70 75 80

Leu Glu Lys Trp Ile Arg Ile Ala Gln Arg Cys Arg Glu Leu Arg Asn 85 90 95

Phe Ser Ser Leu Arg Ala Ile Leu Ser Ala Leu Gln Ser Asn Pro Ile 100 105 110 Tyr Arg Leu Lys Arg Ser Trp Gly Ala Val Ser Arg Glu Pro Leu Ser 115 120 125 Thr Phe Arg Lys Leu Ser Gln Ile Phe Ser Asp Glu Asn Asn His Leu 130 135 140 Ser Ser Arg Glu Ile Leu Phe Gln Glu Glu Ala Thr Glu Gly Ser Gln 145 150 155 160 Glu Glu Asp Asn Thr Pro Gly Ser Leu Pro Ser Lys Pro Pro Pro Gly 165 170 Pro Val Pro Tyr Leu Gly Thr Phe Leu Thr Asp Leu Val Met Leu Asp 180 185 Thr Ala Leu Pro Asp Met Leu Glu Gly Asp Leu Ile Asn Phe Glu Lys 195 200 205 Arg Arg Lys 210 <210> 339 <211> 188 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: RasGEF domain sequence <400> 339 Leu Leu Asp Pro Leu Glu Leu Ala Lys Gln Leu Thr Leu Leu Glu 10

His Glu Leu Phe Lys Lys Ile Asp Pro Phe Glu Cys Leu Gly Gln Val 20 25 30

Trp Gly Lys Lys Tyr Gly Lys Asn Glu Arg Ser Pro Asn Ile Asp Lys
35 40 45

Thr Ile Lys Asn Phe Asn Gln Leu Thr Asn Phe Val Gly Thr Thr Ile 50 55 60

Leu Leu Gln Thr Asp Pro Lys Lys Arg Ala Glu Leu Ile Gln Lys Phe

Ile Gln Val Ala Asp His Cys Arg Glu Leu Asn Asn Phe Asn Ser Leu 85 90 95

Leu Ala Ile Ile Ser Ala Leu Tyr Ser Ser Pro Ile Tyr Arg Leu Lys
100 105 110

Lys Thr Trp Gln Tyr Val Pro Pro Gln Ser Leu Lys Leu Phe Glu Glu
115 120 125

Leu Asn Lys Leu Met Asp Ser Asp Arg Asn Phe Ser Asn Tyr Arg Glu 130 135 140

Leu Leu Lys Ser Ile Phe Pro Leu Pro Cys Val Pro Phe Phe Gly Val
145 150 155 160

Tyr Leu Ser Asp Leu Thr Phe Leu Glu Glu Gly Asn Pro Asp Phe Leu 165 170 175

Glu Thr Asn Leu Val Asn Phe Ser Lys Arg Arg Lys
180 185

<210> 340

<211> 89

<212> PRT

<213> Homo sapiens

<400> 340

Val Leu Arg Val Tyr Phe Gln Asp Leu Lys Pro Gly Val Ala Tyr Lys
1 5 10 15

Thr Ile Arg Val Ser Ser Glu Asp Thr Ala Pro Asp Val Val Gln Leu 20 25 30

Ala Leu Glu Lys Phe Arg Leu Asp Asp Glu Asp Pro Glu Glu Tyr Ala 35 40 45

Leu Val Glu Val Leu Ser Gly Asp Lys Glu Arg Lys Leu Pro Asp Asp 50 55 60

Glu Asn Pro Leu Gln Leu Arg Leu Asn Leu Pro Arg Asp Gly Leu Ser
65 70 75 80

Leu Arg Phe Leu Leu Lys Arg Arg Asp

<210> 341 <211> 89 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Ras association (RalGDS/AF-6) domain sequence <400> 341 Val Leu Arg Val Tyr Phe Gln Asp Leu Lys Pro Gly Val Ala Tyr Lys 5 10 15 1 Thr Ile Arg Val Ser Ser Glu Asp Thr Ala Pro Asp Val Val Gln Leu 20 Ala Leu Glu Lys Phe Arg Leu Asp Asp Glu Asp Pro Glu Glu Tyr Ala 35 40 Leu Val Glu Val Leu Ser Gly Asp Lys Glu Arg Lys Leu Pro Asp Asp 55 60 50 Glu Asn Pro Leu Gln Leu Arg Leu Asn Leu Pro Arg Asp Gly Leu Ser 65 70 75 Leu Arg Phe Leu Leu Lys Arg Arg Asp 85 <210> 342 <211> 83 <212> PRT <213> Homo sapiens <400> 342 Val Ile Arg Val Ser Ile Asp Asn Asp His Gly Asn Leu Tyr Arg Ser 5 10 Ile Leu Leu Thr Ser Gln Asp Lys Ala Pro Ser Val Val Arg Arg Ala 20 25 Leu Gln Lys His Asn Val Pro Gln Pro Trp Ala Cys Asp Tyr Gln Leu

40

55

Phe Gln Val Leu Pro Gly Asp Arg Leu Leu Ile Pro Asp Asn Ala Asn

45

60

35

Arg Lys Glu <210> 343 <211> 86 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Ras association (RalGDS/AF-6) domain sequence Val Leu Arg Val Tyr Phe Asp Asp Pro Gly Gly Thr Tyr Lys Thr Leu 1 5 10 15 Arg Val Ser Lys Arg Thr Thr Ala Arg Asp Val Ile Gln Gln Leu Leu 20 25 30 Glu Lys Phe His Leu Thr Asp Asp Pro Glu Glu Tyr Val Leu Val Glu 35 40 45 Val Lys Glu Gly Gly Lys Glu Arg Val Leu Leu Pro Asp Glu Lys Pro 50 55 60 Leu Gln Leu Gln Lys Leu Trp Pro Arg Gln Gly Ser Asn Leu Arg Phe 65 70 75 80 Val Leu Arg Lys Arg Asp 85 <210> 344 <211> 75 <212> PRT <213> Homo sapiens <400> 344 Asp Pro Ser Phe Met Pro Ala Phe Leu Ala Thr Tyr Arg Thr Phe Val 1 5 10 15 Pro Thr Ala Cys Leu Leu Gly Phe Leu Leu Pro Pro Met Pro Pro Pro 20 25

Val Phe Tyr Ala Met Ser Pro Val Ala Pro Arg Asp Phe Met Leu Arg

75

70

Phe Asn Lys Asn Leu Arg Ala Val Val Ser Val Leu Gly Ser Trp Leu 50 55 60 Gln Asp His Pro Gln Asp Phe Arg Asp Pro Pro 65 70 75 <210> 345 <211> 74 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: RasGEFN domain sequence <400> 345 Asp Pro Thr Phe Val Glu Thr Phe Leu Leu Thr Tyr Arg Ser Phe Ile 15 1 5 10 Thr Thr Gln Glu Leu Leu Gln Lys Leu Leu Tyr Arg Tyr Asn Ala Ile 20 25 30 Pro Pro Glu Gly Val Glu Asp Ile Trp Val Lys Glu Lys Val Asn Pro 35 40 45 Arg Arg Ile Gln Asn Arg Val Leu Asn Ile Leu Arg Leu Trp Val Glu 50 55 60 Asn Tyr Trp Gln Asp Phe Glu Glu Asp Pro 70 65 <210> 346 <211> 184 <212> PRT <213> Homo sapiens <400> 346 Met Ser Arg Leu Ser Arg Ser Leu Leu Trp Ala Ala Thr Cys Leu Gly 1 5 10 15 Val Leu Cys Val Leu Ser Ala Asp Lys Asn Thr Thr Gln His Pro Asn 20 25 30

Pro Pro Pro Gly Val Glu Ile Lys Lys Thr Ala Val Gln Asp Leu Ser

45

40

Ser Leu Pro Leu Val Thr Thr Pro Ala Pro Glu Thr Cys Glu Gly Arg Asn Ser Cys Val Ser Cys Phe Asn Val Ser Val Val Asn Thr Thr Cys Phe Trp Ile Glu Cys Lys Asp Glu Ser Tyr Cys Ser His Asn Ser Thr Val Ser Asp Cys Gln Val Gly Asn Thr Thr Asp Phe Cys Ser Ala Lys Pro Thr Val Gln Pro Ser Pro Ser Thr Thr Ser Lys Thr Val Thr Thr Ser Gly Thr Thr Asn Asn Thr Val Thr Pro Thr Ser Gln Pro Val Arg Lys Ser Thr Phe Asp Ala Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Gly Val Gln Ala Val Ile Phe Phe Leu Tyr Lys Phe Cys Lys Ser Lys Glu Arg Asn Tyr His Thr Leu <210> 347 <211> 197 <212> PRT <213> Homo sapiens <400> 347 Met Ser Arg Leu Ser Arg Ser Leu Leu Trp Ala Ala Thr Cys Leu Gly Val Leu Cys Val Leu Ser Ala Asp Lys Asn Thr Thr Gln His Pro Asn Val Thr Thr Leu Ala Pro Ile Ser Asn Val Thr Ser Ala Pro Val Thr Ser Leu Pro Leu Val Thr Thr Pro Ala Pro Glu Thr Cys Glu Gly Arg

Val Thr Thr Leu Ala Pro Ile Ser Asn Val Thr Ser Ala Pro Val Thr

Asn Ser Cys Val Ser Cys Phe Asn Val Ser Val Val Asn Thr Thr Cys
65 70 75 80

Phe Trp Ile Glu Cys Lys Asp Glu Ser Tyr Cys Ser His Asn Ser Thr 85 90 95

Val Ser Asp Cys Gln Val Gly Asn Thr Thr Asp Phe Cys Ser Val Ser 100 105 110

Thr Ala Thr Pro Val Pro Thr Ala Asn Ser Thr Ala Lys Pro Thr Val 115 120 125

Gln Pro Ser Pro Ser Thr Thr Ser Lys Thr Val Thr Thr Ser Gly Thr 130 135 140

Thr Asn Asn Thr Val Thr Pro Thr Ser Gln Pro Val Arg Lys Ser Thr 145 150 155 160

Phe Asp Ala Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Gly Val 165 170 175

Gln Ala Val Ile Phe Phe Leu Tyr Lys Phe Cys Lys Ser Lys Glu Arg 180 185 190

Asn Tyr His Thr Leu 195

<210> 348

<211> 178

<212> PRT

<213> Homo sapiens

<400> 348

Met Ser Arg Leu Ser Arg Ser Leu Leu Trp Ala Ala Thr Cys Leu Gly
1 5 10 15

Val Leu Cys Val Leu Ser Ala Asp Lys Asn Thr Thr Gln His Pro Asn 20 25 30

Val Thr Thr Leu Ala Pro Ile Ser Asn Val Thr Ser Ala Pro Val Thr 35 40 45

Ser Leu Pro Leu Val Thr Thr Pro Ala Pro Glu Thr Cys Glu Gly Arg
50 60

Asn Ser Cys Val Ser Cys Phe Asn Val Ser Val Val Asn Thr Thr Cys

Phe Trp Ile Glu Cys Lys Asp Glu Ser Tyr Cys Ser His Asn Ser Thr 85 90 95

Val Ser Asp Cys Gln Val Gly Asn Thr Thr Asp Phe Cys Ser Val Ser 100 105 110

Thr Ala Thr Pro Val Pro Thr Ala Asn Ser Thr Gly Thr Thr Asn Asn 115 120 125

Thr Val Thr Pro Thr Ser Gln Pro Val Arg Lys Ser Thr Phe Asp Ala 130 135 140

Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Gly Val Gln Ala Val 145 150 155 160

Ile Phe Phe Leu Tyr Lys Phe Cys Lys Ser Lys Glu Arg Asn Tyr His 165 170 175

Thr Leu

<210> 349

<211> 189

<212> PRT

<213> Homo sapiens

<400> 349

Met Ser Arg Leu Ser Arg Ser Leu Leu Trp Ala Ala Thr Cys Leu Gly
1 5 10 15

Val Leu Cys Val Leu Ser Ala Asp Lys Asn Thr Thr Gln His Pro Asn
20 25 30

Val Thr Thr Leu Ala Pro Ile Ser Asn Val Thr Ser Ala Pro Val Thr 35 40 45

Ser Leu Pro Leu Val Thr Thr Pro Ala Pro Glu Thr Cys Glu Gly Arg
50 55 60

Asn Ser Cys Val Ser Cys Phe Asn Val Ser Val Val Asn Thr Thr Cys 65 70 75 80

Phe Trp Ile Glu Cys Lys Asp Glu Ser Tyr Cys Ser His Asn Ser Thr 85 90 95 Val Ser Asp Cys Gln Val Gly Asn Thr Thr Asp Phe Cys Ser Val Ser Thr Ala Thr Pro Val Pro Thr Ala Asn Ser Thr Ala Lys Pro Thr Val Gln Pro Ser Pro Ser Thr Thr Ser Lys Thr Val Thr Thr Ser Gly Thr Thr Asn Asn Thr Val Thr Pro Thr Ser Gln Pro Val Arg Lys Ser Thr Phe Asp Ala Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Glu Ile Arg Cys His Thr Arg Asn Tyr Ile Pro Asp Leu Lys Lys <210> 350 <211> 195 <212> PRT <213> Rattus norvegicus <400> 350 Met Ser Gly Ala Ser Arg Gly Leu Phe Trp Ala Ala Thr Cys Leu Ala Ala Leu Cys Leu Ser Ala Ala Gln Ser Asn Ser Ser Ala Ser Pro Asn Val Thr Asp Pro Pro Thr Thr Thr Ser Lys Val Val Pro Thr Thr Leu Thr Thr Lys Pro Pro Glu Thr Cys Glu Ser Phe Asn Ser Cys Val Ser Cys Val Asn Ala Thr Leu Thr Asn Asn Ile Thr Cys Val Trp Leu Asp Cys His Glu Ala Asn Lys Thr Tyr Cys Ser Ser Glu Leu Val Ser Asn Cys Thr Gln Lys Thr Ser Thr Asp Ser Cys Ser Val Ile Pro Thr Thr Pro Val Pro Thr Asn Ser Thr Ala Lys Pro Thr Thr Arg Pro Ser

Ser Pro Thr Pro Thr Pro Ser Val Val Thr Ser Ala Gly Ala Thr Asn 130 135 140

Thr Thr Val Thr Pro Thr Ser Gln Pro Glu Arg Lys Ser Thr Phe Asp 145 150 155 160

Ala Ala Ser Phe Ile Gly Gly Ile Val Leu Val Leu Gly Val Gln Ala 165 170 175

Val Ile Phe Phe Leu Tyr Lys Phe Cys Lys Ser Lys Glu Arg Asn Tyr 180 185 190

His Thr Leu 195

<210> 351

<211> 407

<212> PRT

<213> Homo sapiens

<400> 351

Met Ala Val Pro Trp Leu Val Leu Leu Leu Ala Leu Pro Ile Phe Phe 1 5 10 15

Leu Gly Val Phe Val Trp Ala Val Phe Glu His Phe Leu Thr Thr Asp 20 25 30

Ile Pro Ala Thr Leu Gln His Pro Ala Lys Leu Arg Phe Leu His Cys
35 40 45

Ile Phe Leu Tyr Leu Val Thr Leu Gly Asn Ile Phe Glu Lys Leu Gly 50 55 60

Ile Cys Ser Met Pro Lys Phe Ile Arg Phe Leu His Asp Ser Val Arg 65 70 75 80

Ile Lys Lys Asp Pro Glu Leu Val Val Thr Asp Leu Arg Phe Gly Thr
85 90 95

Ile Pro Val Arg Leu Phe Gln Pro Lys Ala Ala Ser Ser Arg Pro Arg 100 105 110

Arg Gly Ile Ile Phe Tyr His Gly Gly Ala Thr Val Phe Gly Ser Leu 115 120 125

Asp Cys Tyr His Gly Leu Cys Asn Tyr Leu Ala Arg Glu Thr Glu Ser

| 130 | 135 | 140 |
|-----|-----|-----|
| | | |

| Val 145 | Leu | Leu | Met | Ile | Gly 150 | Tyr | Arg | Lys | Leu | Pro 155 | Asp | His | His | Ser | Pro 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Leu | Phe | Gln | Asp 165 | Суѕ | Met | Asn | Ala | Ser 170 | Ile | His | Phe | Leu | Lys 175 | Ala |
| Leu | Glu | Thr | Tyr 180 | Gly | Val | Asp | Pro | Ser 185 | Arg | Val | Val | Val | Cys 190 | Gly | Glu |
| Ser | Val | Gly 195 | Gly | Ala | Ala | Val | Ala 200 | Ala | Ile | Thr | Gln | Ala 205 | Leu | Val | Gly |
| Arg | Ser 210 | Asp | Leu | Pro | Arg | Ile 215 | Arg | Ala | Gln | Val | Leu 220 | Ile | Tyr | Pro | Val |
| Val 225 | Gln | Ala | Phe | Cys | Leu 230 | Gln | Leu | Pro | Ser | Phe 235 | Gln | Gln | Asn | Gln | Asn 240 |
| Val | Pro | Leu | Leu | Ser 245 | Arg | Lys | Phe | Met | Val 250 | Thr | Ser | Leu | Cys | Asn 255 | Tyr |
| Leu | Ala | Ile | Asp 260 | Leu | Ser | Trp | Arg | Asp 265 | Ala | Ile | Leu | Asn | Gly 270 | Thr | Cys |
| Val | Pro | Pro 275 | Asp | Val | Trp | Arg | Lys 280 | Tyr | Glu | Lys | Trp | Leu 285 | Ser | Pro | Asp |
| Asn | Ile 290 | Pro | Lys | Lys | Phe | Lys 295 | Asn | Arg | Gly | Tyr | Gln 300 | Pro | Trp | Ser | Pro |
| Gly 305 | Pro | Phe | Asn | Glu | Ala 310 | Ala | Tyr | Leu | Glu | Ala 315 | Lys | His | Met | Leu | Asp 320 |
| Val | Glu | Asn | Ser | Pro 325 | Leu | Ile | Ala | Asp | Asp 330 | Glu | Val | Ile | Ala | Gln 335 | Leu |
| Pro | Glu | Ala | Phe 340 | Leu | Val | Ser | Cys | Glu 345 | Asn | Asp | Ile | Leu | Arg 350 | Asp | Asp |
| Ser | Leu | Leu 355 | Tyr | Lys | Lys | Arg | Leu 360 | Glu | Asp | Gln | Gly | Val 365 | Arg | Val | Thr |
| Trp | Tyr 370 | His | Leu | Tyr | Asp | Gly 375 | Phe | His | Gly | Ser | Ile 380 | Ile | Phe | Phe | Asp |
| Lys | Lys | Ala | Leu | Ser | Phe | Pro | Cys | Ser | Leu | Lys | Ile | Val | Asn | Ala | Val |

385 390 395 400

Val Ser Tyr Ile Lys Gly Ile 405

<210> 352

<211> 409

<212> PRT

<213> Homo sapiens

<400> 352

Met Lys Lys Thr Glu Asp Asn Asn Thr Leu Val Phe Ser Val Asp Val 1 5 10 15

Lys Ala Asn Asn Gly Trp Pro Pro Cys Glu Thr Glu Ser Pro Pro Leu 20 25 30

His Leu Pro Ala Ala Val Asp Met Asp Leu Pro Pro Leu Lys Tyr Asp
35 40 45

Pro Asp Val Val Val Thr Asp Phe Arg Phe Gly Thr Ile Pro Val Lys 50 55 60

Leu Tyr Gln Ser Lys Ala Ser Thr Cys Thr Leu Lys Pro Gly Ile Val 65 70 75 80

Tyr Tyr His Gly Gly Gly Val Met Gly Ser Leu Ser Lys Asn His 85 90 95

Phe Leu Arg Pro Pro Lys Gly Met Asp Trp Arg Val Gly Val Leu Glu
100 105 110

Lys Val Val Gln Ala Val Pro Arg Arg Ile Ser Glu Lys Ile Asp 115 120 125

Arg Lys Phe Ala Gly Val Glu Glu Asn Leu Val Gly Ile Gly Pro Ser 130 135 140

Phe Pro Val Pro Val Arg Asp Cys Leu Val Ala Thr Ile His Phe Leu 165 170 175

Lys Ser Leu Asp Ala Tyr Gly Val Asp Pro Ala Arg Val Val Cys 180 185 190 Gly Asp Ser Phe Gly Gly Ala Ile Ala Ala Val Val Cys Gln Gln Leu 195 200 205

Val Asp Arg Pro Asp Leu Pro Arg Ile Arg Ala Gln Ile Leu Ile Tyr 210 215 220

Ala Ile Leu Gln Ala Leu Asp Leu Gln Thr Pro Ser Phe Gln Gln Arg 225 230 235 240

Lys Asn Ile Pro Leu Leu Thr Trp Ser Phe Ile Cys Tyr Cys Phe Phe 245 250 255

Gln Asn Leu Asp Phe Ser Ser Ser Trp Gln Glu Val Ile Met Lys Gly 260 265 270

Ala His Leu Pro Ala Glu Val Trp Glu Lys Tyr Arg Lys Trp Leu Gly
275 280 285

Pro Glu Asn Ile Pro Glu Arg Phe Lys Glu Arg Gly Tyr Gln Leu Lys 290 . 295 300

Pro His Glu Pro Met Asn Glu Ala Ala Tyr Leu Glu Val Ser Val Val 305 310 315 320

Leu Asp Val Met Cys Ser Pro Leu Ile Ala Glu Asp Asp Ile Val Ser 325 330 335

Gln Leu Pro Glu Thr Cys Ile Val Ser Cys Glu Tyr Asp Ala Leu Arg 340 345 350

Asp Asn Ser Leu Leu Tyr Lys Lys Arg Leu Glu Asp Leu Gly Val Pro 355 360 365

Val Thr Trp His His Met Glu Asp Gly Phe His Gly Val Leu Arg Thr 370 380

Ile Asp Met Ser Phe Leu His Phe Pro Cys Ser Met Arg Ile Leu Ser 385 390 395 400

Ala Leu Val Gln Phe Val Lys Gly Leu 405

<210> 353

<211> 398

<212> PRT

<213> Orycctolagus cuniculus

| V400/ 33 | 00> 353 |
|----------|---------|
|----------|---------|

- Gly Val Lys Thr Val Leu Leu Leu Ile Val Gly Val Leu Gly Ala Tyr
 1 5 10 15
- Tyr Val Tyr Thr Pro Leu Pro Asp Asn Ile Glu Glu Pro Trp Arg Leu 20 25 30
- Leu Trp Val Asn Ala His Met Lys Thr Leu Thr Asn Leu Ala Leu Phe 35 40 45
- Ala Glu Tyr Leu Gly Ser Asn Ile Phe Met Asn Thr Val Lys Phe Leu 50 55 60
- Thr Ser Phe Gln Glu Val Pro Pro Thr Ser Asp Glu Asn Val Thr Val 65 70 75 80
- Thr Glu Thr Thr Phe Asn Asn Val Pro Val Arg Val Tyr Val Pro Lys 85 90 95
- Arg Lys Ser Lys Thr Leu Arg Arg Gly Leu Phe Tyr Ile His Gly Gly
 100 105 110
- Gly Trp Cys Val Gly Ser Ala Ala Leu Ser Gly Tyr Asp Leu Leu Ser 115 120 125
- Arg Arg Thr Ala Asp Arg Leu Asp Val Val Val Ser Thr Asn Tyr 130 135 140
- Asp Ala Leu Lys Trp Phe Leu Arg Gln Asp Val Leu Glu Lys Tyr Gly 165 170 175
- Val Asp Pro Glu Arg Val Gly Val Ser Gly Asp Ser Ala Gly Gly Asn 180 185 190
- Leu Ala Ala Val Ala Gln Gln Leu Ile Lys Asp Pro Asp Val Lys
 195 200 205
- Ile Lys Leu Lys Thr Gln Ser Leu Ile Tyr Pro Ala Leu Gln Thr Leu 210 215 220
- Asp Met Asp Leu Pro Ser Tyr Arg Glu Asn Ala Gln Phe Pro Ile Leu 225 230 235 240
- Ser Lys Ser Phe Met Val Arg Leu Trp Ser Glu Tyr Phe Thr Ser Asp 245 250 255

Arg Ser Leu Glu Lys Ala Met Leu Leu Asn Gln His Val Pro Val Glu 270 260 265 Ser Ser His Leu Phe Lys Phe Thr Asn Trp Ser Ser Leu Leu Pro Glu 280 275 Lys Phe Lys Lys Gly His Val Tyr Asn Thr Pro Thr Tyr Gly Ser Ser 295 Glu Leu Ala Arg Lys Tyr Pro Gly Phe Leu Asp Val Arg Ala Ala Pro 305 310 315 Leu Leu Ala Asp Asp Ala Gln Leu Arg Gly Phe Pro Leu Thr Tyr Val 325 330 Ile Thr Cys Gln Tyr Asp Val Leu Arg Asp Asp Gly Val Met Tyr Val 340 345 Thr Arg Leu Arg Asn Ala Gly Val Gln Val Thr His Asn His Ile Glu 360 365 Asp Gly Phe His Gly Ala Leu Ser Tyr Asn Gly Phe Lys Thr Gly Tyr 375 380 370 Arg Val Glu Lys Gln Tyr Phe Glu Trp Leu Arg Glu Asn Val 390 395 385 <210> 354 <211> 399 <212> PRT <213> Homo sapiens <400> 354 Met Gly Arg Lys Ser Leu Tyr Leu Leu Ile Val Gly Ile Leu Ile Ala 10 Tyr Tyr Ile Tyr Thr Pro Leu Pro Asp Asn Val Glu Glu Pro Trp Arg 25 Met Met Trp Ile Asn Ala His Leu Lys Thr Ile Gln Asn Leu Ala Thr 35 40

Phe Val Glu Leu His Gly Ser Ser Ile Phe Met Asp Ser Phe Lys Val

Val Gly Ser Phe Asp Glu Val Pro Pro Thr Ser Asp Glu Asn Val Thr

| Val | Thr | Glu | Thr | Lys | Phe | Asn | Asn | Ile | Leu | Val | Arg | Val | Tyr | Val | Pro |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | | | 85 | | | | | 90 | | | | | 95 | |

- Lys Arg Lys Ser Glu Ala Leu Arg Arg Gly Leu Phe Tyr Ile His Gly
 100 105 110
- Gly Gly Trp Cys Val Gly Ser Ala Ala Leu Ser Gly Tyr Asp Leu Leu 115 120 125
- Ser Arg Trp Thr Ala Asp Arg Leu Asp Ala Val Val Ser Thr Asn 130 135 140
- Tyr Arg Leu Ala Pro Lys Tyr His Phe Pro Ile Gln Phe Glu Asp Val 145 150 155 160
- Tyr Asn Ala Leu Arg Trp Phe Leu Arg Lys Lys Val Leu Ala Lys Tyr 165 170 175
- Gly Val Asn Pro Glu Arg Ile Gly Ile Ser Gly Asp Ser Ala Gly Gly
 180 185 190
- Asn Leu Ala Ala Ala Val Thr Gln Gln Leu Leu Asp Asp Pro Asp Val 195 200 205
- Lys Ile Lys Leu Lys Ile Gln Ser Leu Ile Tyr Pro Ala Leu Gln Pro 210 215 220
- Leu Asp Val Asp Leu Pro Ser Tyr Gln Glu Asn Ser Asn Phe Leu Phe 225 230 235 240
- Leu Ser Lys Ser Leu Met Val Arg Phe Trp Ser Glu Tyr Phe Thr Thr 245 250 255
- Asp Arg Ser Leu Glu Lys Ala Met Leu Ser Arg Gln His Val Pro Val 260 265 270
- Glu Ser Ser His Leu Phe Lys Phe Ile Asn Trp Ser Ser Leu Leu Pro 275 280 285
- Glu Arg Phe Ile Lys Gly His Val Tyr Asn Asn Pro Asn Tyr Gly Ser 290 295 300
- Ser Glu Leu Ala Lys Lys Tyr Pro Gly Phe Leu Asp Val Arg Ala Ala 305 310 315 320
- Pro Leu Leu Ala Asp Asp Asn Lys Leu Arg Gly Leu Pro Leu Thr Tyr

325 330 335

Val Ile Thr Cys Gln Tyr Asp Leu Leu Arg Asp Asp Gly Leu Met Tyr 340 345 350

Val Thr Arg Leu Arg Asn Thr Gly Val Gln Val Thr His Asn His Val 355 360 365

Glu Asp Gly Phe His Gly Ala Phe Ser Phe Leu Gly Leu Lys Ile Ser 370 375 380

His Arg Leu Ile Asn Gln Tyr Ile Glu Trp Leu Lys Glu Asn Leu 385 390 395

<210> 355

<211> 398

<212> PRT

<213> Rattus norvegicus

<400> 355

Met Gly Arg Thr Ile Phe Leu Leu Ile Ser Val Val Leu Val Ala Tyr 1 5 10 15

Tyr Ile Tyr Ile Pro Leu Pro Asp Asp Ile Glu Glu Pro Trp Lys Ile
20 25 30

Ile Leu Gly Asn Thr Leu Leu Lys Leu Gly Gly Asp Leu Ala Ser Phe 35 40 45

Gly Glu Leu Leu Gly Leu Asn His Phe Met Asp Thr Val Gln Leu Phe
50 55 60

Met Arg Phe Gln Val Val Pro Pro Thr Ser Asp Glu Asn Val Thr Val 65 70 75 80

Met Glu Thr Asp Phe Asn Ser Val Pro Val Arg Ile Tyr Ile Pro Lys 85 90 95

Arg Lys Ser Thr Thr Leu Arg Arg Gly Leu Phe Phe Ile His Gly Gly
100 105 110

Gly Trp Cys Leu Gly Ser Ala Ala Tyr Phe Met Tyr Asp Thr Leu Ser 115 120 125

Arg Arg Thr Ala His Arg Leu Asp Ala Val Val Ser Thr Asp Tyr 130 135 140

| Gly 145 | Leu | Ala | Pro | Lys | Tyr 150 | His | Phe | Pro | Lys | Gln 155 | Phe | Glu | Asp | Val | Tyr 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Ser | Leu | Arg | Trp 165 | Phe | Leu | Gln | Glu | Asp 170 | Ile | Leu | Glu | Lys | Tyr 175 | Gly |
| Val | Asp | Pro | Arg 180 | Arg | Val | Gly | Val | Ser 185 | Gly | Asp | Ser | Ala | Gly 190 | Gly | Asn |
| Leu | Thr | Ala 195 | Ala | Val | Thr | Gln | Gln 200 | Ile | Leu | Gln | Asp | Pro 205 | Asp | Val | Lys |
| Ile | Lys 210 | Leu | Lys | Val | Gln | Ala 215 | Leu | Ile | Tyr | Pro | Ala 220 | Leu | Gln | Ala | Leu |
| Asp 225 | Met | Asn | Val | Pro | Ser 230 | Gln | Gln | Glu | Asn | Ser 235 | Gln | Tyr | Pro | Leu | Leu 240 |
| Thr | Arg | Ser | Leu | Leu 245 | Ile | Arg | Phe | Trp | Ser 250 | Glu | Tyr | Phe | Thr | Thr 255 | Asp |
| Arg | Asp | Leu | Glu 260 | Lys | Ala | Met | Leu | Leu 265 | Asn | Gln | His | Val | Pro 270 | Val | Glu |
| Phe | Ser | His 275 | Leu | Leu | Gln | Phe | Val 280 | Asn | Trp | Ser | Ser | Leu 285 | Leu | Pro | Gln |
| Arg | Tyr 290 | Lys | Lys | Gly | Tyr | Phe 295 | Tyr | Lys | Thr | Pro | Thr 300 | Pro | Gly | Ser | Leu |
| Glu 305 | Leu | Ala | Gln | Lys | Tyr 310 | Pro | Gly | Phe | Thr | Asp 315 | Val | Lys | Ala | Cys | Pro 320 |
| Leu | Leu | Ala | Asn | Asp 325 | Ser | Ile | Leu | His | His 330 | Leu | Pro | Met | Thr | Tyr 335 | Ile |
| Ile | Thr | Cys | Gln 340 | Tyr | Asp | Val | Leu | Arg 345 | Asp | Asp | Gly | Leu | Met 350 | Tyr | Val |
| Lys | Arg | Leu 355 | Gln | Asn | Thr | Gly | Val 360 | His | Val | Thr | His | His 365 | His | Ile | Glu |
| Asp | Gly 370 | Phe | His | Gly | Ala | Leu 375 | Thr | Leu | Pro | Gly | Leu 380 | Lys | Ile | Thr | Tyr |
| Arg 385 | Met | Gln | Asn | Gln | Tyr 390 | Leu | Asn | Trp | Leu | His 395 | Lys | Asn | Leu | | |

<211> 109 <212> PRT <213> Homalozoon vermiculare <400> 356 Leu Phe Gln Pro Lys Ala Ala Ser Ser Arg Pro Arg Arg Gly Ile Ile Phe Tyr His Gly Gly Ala Thr Val Phe Gly Ser Leu Asp Cys Tyr His 20 25 Gly Leu Cys Asn Tyr Leu Ala Arg Glu Thr Glu Ser Val Leu Leu Met 40 Ile Gly Tyr Arg Lys Leu Pro Asp His His Ser Pro Ala Leu Phe Gln Asp Cys Met Asn Ala Ser Ile His Phe Leu Lys Ala Leu Glu Thr Tyr 75 70 Gly Val Asp Pro Ser Arg Val Val Cys Gly Glu Ser Val Gly Gly Ala Ala Val Ala Ala Ile Thr Gln Ala Leu Val Gly Arg 100 105 <210> 357 <211> 118 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Carboxylesterase domain sequence <400> 357 Val Tyr Thr Pro Lys Asn Arg Lys Pro Asn Ser Lys Leu Pro Val Met 1 5 10 15 Val Trp Ile His Gly Gly Gly Phe Met Phe Gly Ser Gly Leu Ser Leu 20 25 30 Tyr Asp Gly Glu Ser Leu Ala Arg Glu Gly Asn Val Ile Val Val Ser 35

<210> 356

40

Ile Asn Tyr Arg Leu Gly Pro Leu Gly Phe Leu Ser Thr Gly Asp Asp Val Leu Pro Gly Asn Tyr Gly Leu Leu Asp Gln Arg Leu Ala Leu Lys Trp Val Gln Asp Asn Ile Ala Ala Phe Gly Gly Asp Pro Asp Ser Val Thr Ile Phe Gly Glu Ser Ala Gly Gly Ala Ser Val Ser Leu Leu Leu Leu Ser Pro Ser Ser Lys <210> 358 <211> 280 <212> PRT <213> Homo sapiens <400> 358 Met Leu Leu Gly Asn Leu Ala Ile Ile Ser Phe Ile Cys Leu Asp Ser Arg Leu His Ser Pro Met Tyr Phe Phe Leu Cys Asn Phe Ser Leu Met Glu Met Val Val Thr Ser Thr Val Val His Arg Met Leu Ala Asp Leu Leu Ser Thr His Lys Thr Met Ser Leu Ala Lys Cys Leu Thr Gln Ser Phe Phe Tyr Phe Ser Leu Gly Ser Ala Asn Phe Leu Ile Leu Met Val Met Ala Phe Asp Arg Tyr Val Ala Ile Cys His Pro Leu Arg Tyr Pro Thr Ile Thr Asn Gly Pro Val Cys Val Lys Leu Val Val Ala Cys Trp Val Val Gly Phe Leu Ser Ile Val Ser Pro Thr Leu Gln Lys Thr Arg Leu Trp Phe Cys Gly Pro Asn Ile Ile Gly His Tyr Phe Cys Asp Ser

Ala Pro Leu Lys Leu Ala Cys Ser Asp Thr Arg His Ile Glu Arg Met Asp Leu Phe Leu Ser Leu Leu Phe Val Leu Thr Thr Met Leu Leu Ile Ile Leu Ser Tyr Ile Leu Ile Val Ala Ala Val Leu His Ile Pro Ser Ser Ser Gly Cys Gln Lys Ala Phe Ser Thr Cys Ala Ser His Leu Thr Val Val Leu Gly Tyr Gly Ser Ala Ile Phe Ile Tyr Val Arg Pro Gly Lys Gly His Ser Thr Tyr Leu Asn Lys Ala Val Ala Met Val Thr Ala Met Val Thr Pro Phe Leu Asn Pro Phe Ile Phe Thr Phe Arg Asn Glu Lys Val Lys Glu Val Ile Glu Asp Val Thr Lys Arg Ile Phe Leu Gly Asp Pro Ala Ala Cys Arg <210> 359 <211> 216 <212> PRT <213> Homo sapiens <400> 359 Leu Met Glu Met Val Val Thr Ser Thr Val Val His Arg Met Leu Ala Asp Leu Leu Ser Thr His Lys Thr Met Ser Leu Ala Lys Cys Leu Thr Gln Ser Phe Phe Tyr Phe Ser Leu Gly Ser Ala Asn Phe Leu Ile Leu Met Val Met Ala Phe Asp Arg Tyr Val Ala Ile Cys His Pro Leu Arg

Tyr Pro Thr Ile Thr Asn Gly Pro Val Cys Val Lys Leu Val Val Ala

Cys Trp Val Val Gly Phe Leu Ser Ile Val Ser Pro Thr Leu Gln Lys 85 90 95

Thr Arg Leu Trp Phe Cys Gly Pro Asn Ile Ile Gly His Tyr Phe Cys 100 105 110

Asp Ser Ala Pro Leu Leu Lys Leu Ala Cys Ser Asp Thr Arg His Ile 115 120 125

Glu Arg Met Asp Leu Phe Leu Ser Leu Leu Phe Val Leu Thr Thr Met 130 135 140

Leu Leu Ile Ile Leu Ser Tyr Ile Leu Ile Val Ala Ala Val Leu His 145 150 155 160

Ile Pro Ser Ser Ser Gly Cys Gln Lys Ala Phe Ser Thr Cys Ala Ser 165 170 175

His Leu Thr Val Val Leu Gly Tyr Gly Ser Ala Ile Phe Ile Tyr 180 185 190

Val Arg Pro Gly Lys Gly His Ser Thr Tyr Leu Asn Lys Ala Val Ala 195 200 205

Met Val Thr Ala Met Val Thr Pro 210 215

<210> 360

65

<211> 1056

<212> PRT

<213> Homo sapiens

<400> 360

Met Pro Val Leu Leu Pro Val His Phe Ser Ala Lys Cys Pro Leu Leu
1 5 10 15

Leu Leu Cys Asp Pro Ala Asn Pro Pro Ser Glu Pro Leu Pro Ser Gln 20 25 30

Gly Cys Phe Ile Phe Ile His Arg Val Leu Leu Asp Leu Ser Thr Ala $35 \hspace{1.5cm} 40 \hspace{1.5cm} 45$

Gly Glu Ser Gly Asn Thr Ala Gly Phe Ile Cys Asp Gln Ala Leu Leu 50 55 60

| Thr 65 | Ser | Pro | Val | Arg | Glu 70 | Asp | Gly | Ala | Glu | Asn 75 | Gly | Leu | Gly | Phe | His 80 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gln | Pro | Val | Glu | Leu 85 | His | Ile | Cys | Gly | Asp 90 | Ala | Val | Gly | Phe | Val 95 | Gly |
| Met | Gly | Gln | Arg 100 | Arg | Lys | Pro | Met | Ser 105 | Val | Pro | Trp | Ser | His 110 | Pro | Lys |
| Ile | Ser | Glu 115 | Lys | Cys | Ala | Ser | Asp 120 | Thr | Trp | Cys | Thr | Asp 125 | Ala | Thr | Tyr |
| His | Arg 130 | Glu | His | Ser | Lys | Pro 135 | Ser | Gly | Pro | Trp | Glu 140 | His | Gly | Pro | Leu |
| Lys 145 | Pro | Phe | Glu | Asp | Trp 150 | Val | Pro | Ala | Leu | Pro 155 | Tyr | Pro | Leu | Trp | Pro 160 |
| Gln | Glu | Leu | Leu | His 165 | Cys | Gly | Ser | Gln | Ser 170 | Gly | Asp | Cys | Met | Cys 175 | Leu |
| Leu | Leu | Leu | Glu 180 | Ser | Ser | Arg | Arg | Ser 185 | Pro | Pro | Thr | Leu | Pro 190 | Ile | Pro |
| Leu | Thr | Phe 195 | Pro | Arg | Leu | Cys | Gln 200 | Ser | Phe | Pro | Leu | Leu 205 | Thr | Ala | Ser |
| Gly | Lys 210 | Glu | Pro | Ser | Cys | Gly 215 | Phe | Thr | Ser | Ala | Leu 220 | Arg | Arg | Leu | Tyr |
| Gly 225 | Cys | Gly | Ala | Ala | Glu 230 | Arg | Pro | Gln | Ser | Pro 235 | Val | Thr | Pro | Lys | Thr 240 |
| Glu | Thr | Ser | Glu | Gln 245 | Gly | Pro | Lys | Asp | Pro 250 | Pro | Ile | His | Leu | Ala 255 | His |
| Pro | Ser | Asp | Arg 260 | Ala | Leu | Ser | Pro | Ser 265 | Cys | Phe | Leu | Ser | Leu 270 | Arg | Ala |
| Val | Ile | Leu 275 | Thr | Cys | Lys | Asn | Arg 280 | Asp | Ala | Gln | Val | Glu 285 | Glu | Gly | His |
| Arg | Arg 290 | Glu | Pro | Pro | Val | Leu 295 | Asp | Cys | Gly | Tyr | Gln 300 | Arg | Ser | Gly | Thr |
| Arg 305 | Gly | Asn | His | Thr | Arg 310 | Arg | Ile | Cys | Ser | Thr 315 | Leu | Arg | Gly | Ser | Arg 320 |

| Ile | Glu | Ala | Trp | Val 325 | Ala | Ala | Ala | Thr | Leu 330 | Gln | Arg | Gly | Pro | Tyr 335 | Phe |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Arg | Lys | Gln | Gln 340 | Pro | Leu | Gly | Lys | Asp 345 | Ser | Trp | Ser | Val | Ala 350 | Glu | Asp |
| Trp | Ile | Glu 355 | Ala | Phe | Met | Leu | Ala 360 | Phe | Gly | Val | Arg | Val 365 | Leu | Trp | Asp |
| Ala | Ser 370 | Met | Ala | Leu | Glu | Ala 375 | Gln | Arg | Asp | Pro | Ser 380 | Ser | Asn | Asp | Thr |
| Lys 385 | Gly | Lys | Asp | Gln | Leu 390 | Thr | Lys | Arg | Asp | Gln 395 | Arg | Asn | Pro | Gln | Asn 400 |
| Phe | Ala | Leu | Leu | Gln 405 | Lys | Ser | Ala | Ala | Ser 410 | Asp | Trp | Asn | Ser | Gln 415 | Pro |
| Val | Cys | Arg | Arg 420 | Gly | Tyr | Leu | Thr | Cys 425 | Ala | Ser | Ala | Ser | Leu 430 | Gly | Glu |
| Ile | Ser | Ser 435 | Pro | His | Phe | Pro | Val 440 | His | Leu | Asn | Ala | Pro 445 | Lys | Cys | His |
| Trp | Gly 450 | Leu | Ser | Ser | Ser | Pro 455 | Val | Glu | Arg | Trp | Met 460 | Leu | Arg | Glu | Arg |
| Lys 465 | Ala | Val | Thr | Asp | Glu 470 | Ser | Ser | Ser | Ser | Trp 475 | Met | Val | Ala | Ile | Arg 480 |
| Ala | Arg | Glu | Thr | Pro 485 | Gly | Ile | Leu | Ala | Gln 490 | Arg | Ile | Cys | Ser | Ala 495 | Leu |
| Lys | Gly | Val | Trp 500 | Cys | Gln | Ala | Ala | Gln 505 | Gly | Ser | Leu | Pro | Arg 510 | Leu | Leu |
| Ser | Ser | Leu 515 | Ser | Ile | Ser | Thr | Gly 520 | Cys | Asp | Lys | Thr | Ala 525 | Val | Leu | Thr |
| Phe | Asp 530 | Arg | Ala | Leu | Leu | Thr 535 | Arg | Glu | His | Ser | Lys 540 | Pro | Asn | Gly | Pro |
| Trp 545 | Glu | Arg | Gly | Pro | Leu 550 | Lys | Pro | Ser | Gly | Asp 555 | Trp | Asp | Thr | Cys | Leu 560 |
| His | Tyr | Leu | Leu | Trp 565 | Pro | Gln | Glu | Leu | Phe 570 | His | Cys | Arg | Ser | Gln 575 | Thr |

Glu Asp Tyr Thr Val Thr Trp Phe Asp Val Val Asp Arg Gln Met Gln Lys Tyr Ser Gln Ser Pro Phe Leu Glu Gln Arg Val Lys Lys Thr Met Ser Pro Asp Gly Asn His Ser Ser Asp Pro Thr Glu Phe Val Leu Ala Gly Leu Pro Asn Leu Asn Ser Ala Arg Val Glu Leu Phe Ser Val Phe Leu Leu Val Tyr Leu Leu Asn Leu Thr Gly Asn Val Leu Ile Val Gly Val Val Arg Ala Asp Thr Arg Leu Gln Thr Pro Met Tyr Phe Phe Leu Gly Asn Leu Ser Cys Leu Glu Ile Leu Leu Thr Ser Val Ile Ile Pro Lys Met Leu Ser Asn Phe Leu Ser Arg Gln His Thr Ile Ser Phe Ala Ala Cys Ile Thr Gln Phe Tyr Phe Tyr Phe Phe Leu Gly Ala Ser Glu Phe Leu Leu Ala Val Met Ser Ala Asp Arg Tyr Leu Ala Ile Cys His Pro Leu Arg Tyr Pro Leu Leu Met Ser Gly Ala Val Cys Phe Arg Val Ala Leu Ala Cys Trp Val Gly Gly Leu Val Pro Val Leu Gly Pro Thr Val Ala Val Ala Leu Leu Pro Phe Cys Lys Gln Gly Ala Val Val Gln His Phe Phe Cys Asp Ser Gly Pro Leu Leu Arg Leu Ala Cys Thr Asn Thr Lys Lys Leu Glu Glu Thr Asp Phe Val Leu Ala Ser Leu Val Ile Val Ser Ser Leu Leu Ile Thr Ala Val Ser Tyr Gly Leu Ile Val

Leu Ala Val Leu Ser Ile Pro Ser Ala Ser Gly Arg Gln Lys Ala Phe 835 840 845

Ser Thr Cys Thr Ser His Leu Ile Val Val Thr Leu Phe Tyr Gly Ser 850 855 860

Ala Ile Phe Leu Tyr Val Arg Pro Ser Gln Ser Gly Ser Val Asp Thr 865 870 . 875 880

Asn Trp Ala Val Thr Val Ile Thr Thr Phe Val Thr Pro Leu Leu Asn 885 890 895

Pro Phe Ile Tyr Ala Leu Arg Asn Glu Gln Val Lys Glu Ala Leu Lys 900 905 910

Asp Met Phe Arg Lys Gly Cys Asp Phe Ala Phe Glu Arg Cys Asn Ser 915 920 925

Ala Cys Asn Cys Arg Lys Gly Ser Leu Thr Thr Thr Thr Lys Ser Ala 930 935 940

Thr Leu Arg Cys Gly Ala Gly Ala Lys Ala Arg Ala Gly Ala Arg Leu 945 950 955 960

His Pro Ala Ala Gly Ser Pro Arg Asp Ser Arg Lys Val Asn Val Arg 965 970 975

Val Gln Lys Asp Pro Arg Arg Ser Val Pro Lys Val Glu Thr Phe Ile 980 985 990

Ser Gly Ser Gly Pro Ser Cys Val Gly Gln Cys Thr Gly Arg Val Cys 995 1000 1005

Ile Leu Lys Gly Thr Arg Thr Ile Ser Gly Gly Leu Trp Leu Glu Asp 1010 1015 1020

Pro Arg Lys Thr Arg Thr Thr Asp Phe Thr His Arg Lys Ile Lys Val 1025 1030 1035 1040

Thr Ala Gly Leu Ala Gly Glu Lys Val Glu Pro Thr Leu Pro Arg Cys 1045 1050 1055 <212> PRT

<213> Homo sapiens

<400> 361

Met Ala Asn Leu Ser Gln Pro Ser Glu Phe Val Leu Leu Gly Phe Ser 1 5 10 15

Ser Phe Gly Glu Leu Gln Ala Leu Leu Tyr Gly Pro Phe Leu Met Leu 20 25 30

Tyr Leu Leu Ala Phe Met Gly Asn Thr Ile Ile Ile Val Met Val Ile 35 40 45

Ala Asp Thr His Leu His Thr Pro Met Tyr Phe Phe Leu Gly Asn Phe 50 55 60

Ser Leu Leu Glu Ile Leu Val Thr Met Thr Ala Val Pro Arg Met Leu 65 70 75 80

Ser Asp Leu Leu Val Pro His Lys Val Ile Thr Phe Thr Gly Cys Met 85 90 95

Val Gln Phe Tyr Phe His Phe Ser Leu Gly Ser Thr Ser Phe Leu Ile 100 105 110

Leu Thr Asp Met Ala Leu Asp Arg Phe Val Ala Ile Cys His Pro Leu 115 120 125

Arg Tyr Gly Thr Leu Met Ser Arg Ala Met Cys Val Gln Leu Ala Gly 130 135 140

Ala Ala Trp Ala Ala Pro Phe Leu Ala Met Val Pro Thr Val Leu Ser 145 150 155 160

Arg Ala His Leu Asp Tyr Cys His Gly Asp Val Ile Asn His Phe Phe 165 170 175

Cys Asp Asn Glu Pro Leu Leu Gln Leu Ser Cys Ser Asp Thr Arg Leu
180 185 190

Leu Glu Phe Trp Asp Phe Leu Met Ala Leu Thr Phe Val Leu Ser Ser 195 200 205

Phe Leu Val Thr Leu Ile Ser Tyr Gly Tyr Ile Val Thr Thr Val Leu 210 215 220

Arg Ile Pro Ser Ala Ser Ser Cys Gln Lys Ala Phe Ser Thr Cys Gly 225 230 235 240

Ser His Leu Thr Leu Val Phe Ile Gly Tyr Ser Ser Thr Ile Phe Leu 245 250 255

Tyr Val Arg Pro Gly Lys Ala His Ser Val Gln Val Arg Lys Val Val 260 265 270

Ala Leu Val Thr Ser Val Leu Thr Pro Phe Leu Asn Pro Phe Ile Leu 275 280 285

Thr Phe Cys Asn Gln Thr Val Lys Thr Val Leu Gln Gly Gln Met Gln 290 295 300

Arg Leu Lys Gly Leu Cys Lys Ala Gln 305 310

<210> 362

<211> 347

<212> PRT

<213> Homo sapiens

<400> 362

Met Gly Asn Trp Thr Ala Ala Val Thr Glu Phe Val Leu Leu Gly Phe
1 5 10 15

Ser Leu Ser Arg Glu Val Glu Leu Leu Leu Leu Val Leu Leu Leu Pro 20 25 30

Thr Phe Leu Leu Thr Leu Leu Gly Asn Leu Leu Ile Ile Ser Thr Val 35 40 45

Leu Ser Cys Ser Arg Leu His Thr Pro Met Tyr Phe Phe Leu Cys Asn 50 55 60

Leu Ser Ile Leu Asp Ile Leu Phe Thr Ser Val Ile Ser Pro Lys Val 65 70 75 80

Leu Ala Asn Leu Gly Ser Arg Asp Lys Thr Ile Ser Phe Ala Gly Cys 85 90 95

Ile Thr Gln Cys Tyr Phe Tyr Phe Phe Leu Gly Thr Val Glu Phe Leu 100 105 110

Leu Leu Thr Val Met Ser Tyr Asp Arg Tyr Ala Thr Ile Cys Cys Pro 115 120 125

Leu Arg Tyr Thr Thr Ile Met Arg Pro Ser Val Cys Ile Gly Thr Val

| Val 145 | Phe | Ser | Trp | Val | Gly 150 | Gly | Phe | Leu | Ser | Val 155 | Leu | Phe | Pro | Thr | Ile 160 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Ile | Ser | Gln | Leu 165 | Pro | Phe | Cys | Gly | Ser 170 | Asn | Ile | Ile | Asn | His 175 | Phe |
| Phe | Cys | Asp | Ser 180 | Gly | Pro | Leu | Leu | Ala 185 | Leu | Ala | Cys | Ala | Asp 190 | Thr | Thr |
| Ala | Ile | Glu 195 | Leu | Met | Asp | Phe | Met 200 | Leu | Ser | Ser | Met | Val 205 | Ile | Leu | Cys |
| Cys | Ile 210 | Val | Leu | Val | Ala | Tyr 215 | Ser | Tyr | Thr | Tyr | Ile 220 | Ile | Leu | Thr | Ile |
| Val 225 | Arg | Ile | Pro | Ser | Ala 230 | Ser | Gly | Arg | Lys | Lys 235 | Ala | Phe | Asn | Thr | Cys 240 |
| Ala | Ser | His | Leu | Thr 245 | Ile | Val | Ile | Ile | Pro 250 | Ser | Gly | Ile | Thr | Val 255 | Phe |
| Ile | Tyr | Val | Thr 260 | Pro | Ser | Gln | Lys | Glu 265 | Tyr | Leu | Glu | Ile | Asn 270 | Lys | Ile |
| Pro | Leu | Val 275 | Leu | Ser | Ser | Val | Val 280 | Thr | Pro | Phe | Leu | Asn 285 | Pro | Phe | Ile |
| Tyr | Thr 290 | Leu | Arg | Asn | Asp | Thr 295 | Val | Gln | Gly | Val | Leu 300 | Arg | Asp | Val | Trp |
| Val 305 | Arg | Val | Arg | Gly | Val 310 | Phe | Glu | Lys | Arg | Met 315 | Arg | Ala | Val | Leu | Arg 320 |
| Ser | Arg | Leu | Ser | Ser 325 | Asn | Lys | Asp | His | Gln 330 | Gly | Arg | Ala | Cys | Ser 335 | Ser |

<210> 363

<211> 246

<212> PRT

<213> Homo sapiens

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345

Pro Pro Cys Val Tyr Ser Val Lys Leu Gln Cys

| Ile 1 | Ile | Ser | Phe | Ile 5 | Cys | Leu | Asp | Ser | Arg 10 | Leu | His | Ser | Pro | Met 15 | Tyr |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Phe | Phe | Leu | Cys 20 | Asn | Phe | Ser | Leu | Met 25 | Glu | Met | Val | Val | Thr 30 | Ser | Thr |
| Val | Val | His 35 | Arg | Met | Leu | Ala | Asp 40 | Leu | Leu | Ser | Thr | His 45 | Lys | Thr | Met |
| Ser | Leu 50 | Ala | Lys | Cys | Leu | Thr 55 | Gln | Ser | Phe | Phe | Tyr 60 | Phe | Ser | Leu | Gly |
| Ser 65 | Ala | Asn | Phe | Leu | Ile 70 | Leu | Met | Val | Met | Ala 75 | Phe | Asp | Arg | Tyr | Val 80 |
| Ala | Ile | Cys | His | Pro 85 | Leu | Arg | Tyr | Pro | Thr 90 | Ile | Thr | Asn | Gly | Pro 95 | Val |
| Cys | Val | Lys | Leu 100 | Val | Val | Ala | Cys | Trp 105 | Val | Val | Gly | Phe | Leu 110 | Ser | Ile |
| Val | Ser | Pro 115 | Thr | Leu | Gln | Lys | Thr 120 | Arg | Leu | Trp | Phe | Cys 125 | Gly | Pro | Asn |
| Ile | Ile 130 | Gly | His | Tyr | Phe | Cys 135 | Asp | Ser | Ala | Pro | Leu 140 | Leu | Lys | Leu | Ala |
| Cys 145 | Ser | Asp | Thr | Arg | His 150 | Ile | Glu | Arg | Met | Asp 155 | Leu | Phe | Leu | Ser | Leu 160 |
| Leu | Phe | Val | Leu | Thr 165 | Thr | Met | Leu | Leu | Ile 170 | Ile | Leu | Ser | Tyr | Ile 175 | Leu |
| Ile | Val | Ala | Ala 180 | Val | Leu | His | Ile | Pro 185 | Ser | Ser | Ser | Gly | Cys 190 | Gln | Lys |
| Ala | Phe | Ser 195 | Thr | Cys | Ala | Pro | His 200 | Leu | Thr | Val | Val | Val 205 | Leu | Gly | Tyr |
| Gly | Ser 210 | Ala | Ile | Phe | Ile | Tyr 215 | Val | Arg | Pro | Gly | Lys 220 | Gly | His | Ser | Thr |
| Tyr 225 | Leu | Asn | Lys | Ala | Val 230 | Ala | Met | Val | Thr | Ala 235 | Met | Val | Thr | Pro | Phe 240 |
| Leu | Asn | Pro | Phe | Ile | Phe | | | | | | | | | | |

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         35
     50
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195 200 205

Pro Tyr His Ile Val Leu Leu Leu Asp Ser Leu Cys Leu Leu Ser Ile 210 215 220

Trp Arg Val Leu Pro Thr Ala Leu Leu Ile Thr Leu Trp Leu Ala Tyr 225 230 235 240

Val Asn Ser Cys Leu Asn Pro Ile Ile Tyr 245 250

<210> 365

<211> 559

<212> PRT

<213> Homo sapiens

<400> 365

Met Ala Pro Thr Leu Gln Gln Ala Tyr Arg Arg Arg Trp Trp Met Ala
1 5 10 15

Cys Thr Ala Val Leu Glu Asn Leu Phe Phe Ser Ala Val Leu Leu Gly
20 25 30

Trp Gly Ser Leu Leu Ile Ile Leu Lys Asn Glu Gly Phe Tyr Ser Ser 35 40 45

Thr Cys Pro Ala Glu Ser Ser Thr Asn Thr Thr Gln Asp Glu Gln Arg 50 55 60

Arg Trp Pro Gly Cys Asp Gln Gln Asp Glu Met Leu Asn Leu Gly Phe 65 70 75 80

Thr Ile Gly Ser Phe Val Leu Ser Ala Thr Thr Leu Pro Leu Gly Ile 85 90 95

Leu Met Asp Arg Phe Gly Pro Arg Pro Val Arg Leu Val Gly Ser Ala
100 105 110

Cys Phe Thr Ala Ser Cys Thr Leu Met Ala Leu Ala Ser Arg Asp Val 115 120 125

Glu Ala Leu Ser Pro Leu Ile Phe Leu Ala Leu Ser Leu Asn Gly Phe 130 135 140

Gly Gly Ile Cys Leu Thr Phe Thr Ser Leu Thr Leu Pro Asn Met Phe 145 150 155 160

| Gly | Asn | Leu | Arg | Ser 165 | Thr | Leu | Met | Ala | Leu 170 | Met | Ile | Gly | Ser | Tyr 175 | Ala |
|--------------------------|---------------------|---------------------------------|--------------------------|--------------------------|--------------------------|---------------------------------|---------------------------------|---------------------------------|--------------------------|---------------------------------|---------------------------------|--------------------------|--------------------------|---------------------------------|--------------------------|
| Ser | Ser | Ala | Ile 180 | Thr | Phe | Pro | Gly | Ile 185 | Lys | Leu | Ile | Tyr | Asp 190 | Ala | Gly |
| Val | Ala | Phe 195 | Val | Val | Ile | Met | Phe 200 | Thr | Trp | Ser | Gly | Leu 205 | Ala | Cys | Leu |
| Ile | Phe 210 | Leu | Asn | Cys | Thr | Leu 215 | Asn | Trp | Pro | Ile | Glu 220 | Ala | Phe | Pro | Ala |
| Pro 225 | Glu | Glu | Val | Asn | Tyr 230 | Thr | Lys | Lys | Ile | Lys 235 | Leu | Ser | Gly | Leu | Ala 240 |
| Leu | Asp | His | Lys | Val 245 | Thr | Gly | Asp | Leu | Phe 250 | Tyr | Thr | His | Val | Thr 255 | Thr |
| Met | Gly | Gln | Arg 260 | Leu | Ser | Gln | Lys | Ala 265 | Pro | Ser | Leu | Glu | Asp 270 | Gly | Ser |
| Asp | Ala | Phe 275 | Met | Ser | Pro | Gln | Asp 280 | Val | Arg | Gly | Thr | Ser 285 | Glu | Asn | Leu |
| Pro | Glu 290 | Arg | Ser | Val | Pro | Leu 295 | Arg | Lys | Ser | Leu | Cys 300 | Ser | Pro | Thr | Phe |
| | | | | | | | | | | | | | | | |
| Leu 305 | Trp | Ser | Leu | Leu | Thr 310 | Met | Gly | Met | Thr | Gln 315 | Leu | Arg | Ile | Ile | Phe 320 |
| 305 | | | Leu Ala | | 310 | | | | | 315 | | | | | 320 |
| 305 Tyr | Met | Ala | | Val 325 | 310 Asn | Lys | Met | Leu | Glu 330 | 315 Tyr | Leu | Val | Thr | Gly 335 | 320 Gly |
| 305 Tyr Gln | Met Glu | Ala His | Ala Glu | Val 325 Thr | 310 Asn Asn | Lys Glu | Met | Leu Gln 345 | Glu 330 Gln | 315 Tyr Lys | Leu Val | Val Ala | Thr Glu 350 | Gly 335 Thr | 320 Gly Val |
| 305 Tyr Gln Gly | Met Glu Phe | Ala His Tyr 355 | Ala Glu 340 | Val 325 Thr | 310 Asn Asn Val | Lys Glu Phe | Met Gln Gly 360 | Leu Gln 345 Ala | Glu 330 Gln Met | 315 Tyr Lys Gln | Leu Val Leu | Val Ala Leu 365 | Thr Glu 350 Cys | Gly 335 Thr | 320 Gly Val Leu |
| 305 Tyr Gln Gly | Met Glu Phe Cys 370 | Ala His Tyr 355 Pro | Ala Glu 340 Ser | Val 325 Thr Ser | 310 Asn Asn Val | Lys Glu Phe Tyr 375 | Met Gln Gly 360 Ile | Leu Gln 345 Ala Met | Glu 330 Gln Met | 315 Tyr Lys Gln Trp | Leu Val Leu Arg 380 | Val Ala Leu 365 | Thr Glu 350 Cys | Gly 335 Thr Leu Asp | 320 Gly Val Leu Cys |

Thr Asn Ala Ile Ser Ala Phe Thr Leu Thr Asn Leu Leu Leu Val Gly Phe Gly Ile Thr Cys Leu Ile Asn Asn Leu His Leu Gln Phe Val Thr Phe Val Leu His Thr Ile Val Arg Gly Phe Phe His Ser Ala Cys Gly Ser Leu Tyr Ala Ala Val Phe Pro Ser Asn His Phe Gly Thr Leu Thr Gly Leu Gln Ser Leu Ile Ser Ala Val Phe Ala Leu Leu Gln Gln Pro Leu Phe Met Ala Met Val Gly Pro Leu Lys Gly Glu Pro Phe Trp Val Asn Leu Gly Leu Leu Phe Ser Leu Leu Gly Phe Leu Leu Pro Ser Tyr Leu Phe Tyr Tyr Arg Ala Arg Leu Gln Gln Glu Tyr Ala Ala Asn Gly Met Gly Pro Leu Lys Val Leu Ser Gly Ser Glu Val Thr Ala <210> 366 <211> 654 <212> PRT <213> Mus musculus <400> 366 Met Pro Trp Leu Pro Gly Phe Thr Tyr Leu Trp Arg Gln Asp Gly Ser Gln Ile His Cys Phe Phe Arg Gly Arg Arg Gly Glu Thr Gly Gly Ser Glu Ala Arg Trp Val Trp His Ala Gly Lys Thr Pro Arg Val Asp Ala Ile Trp Asn Trp Asp Pro Gly Ser Gln Glu Ile Arg Ser Val Glu Ala Pro Gly Arg Leu Cys Val Thr Pro Gly Val Lys Ser Cys Gly Arg

| Gln | Val | Cys | Arg | Gly 85 | Gln | Ser | Leu | Gly | His 90 | His | Gly | Ser | His | Ala 95 | Glu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Gly | Val | Pro 100 | Gln | Arg | Trp | Trp | Met 105 | Ala | Cys | Thr | Ala | Val 110 | Val | Glu |
| Asn | Leu | Phe 115 | Phe | Ser | Ala | Val | Leu 120 | Leu | Gly | Trp | Ala | Ser 125 | Leu | Leu | Ile |
| Met | Leu 130 | Lys | Lys | Glu | Gly | Phe 135 | Tyr | Ser | Ser | Leu | Cys 140 | Pro | Ala | Glu | Asn |
| Arg 145 | Thr | Asn | Thr | Thr | Gln 150 | Asp | Glu | Gln | His | Gln 155 | Trp | Thr | Ser | Cys | Asp 160 |
| Gln | Gln | Glu | Lys | Met 165 | Leu | Asn | Leu | Gly | Phe 170 | Thr | Ile | Gly | Ser | Phe 175 | Leu |
| Leu | Ser | Ala | Thr 180 | Thr | Leu | Pro | Leu | Gly 185 | Ile | Leu | Met | Asp | Arg 190 | Phe | Gly |
| Pro | Arg | Pro 195 | Leu | Arg | Leu | Val | Gly 200 | Ser | Ala | Cys | Phe | Ala 205 | Ala | Ser | Cys |
| Thr | Leu 210 | Met | Ala | Leu | Ala | Ser 215 | Arg | Asp | Thr | Glu | Val 220 | Leu | Ser | Pro | Leu |
| Ile 225 | Phe | Leu | Ala | Leu | Ser 230 | Leu | Asn | Gly | Phe | Ala 235 | Gly | Ile | Cys | Leu | Thr 240 |
| Phe | Thr | Ser | Leu | Thr 245 | Leu | Pro | Asn | Met | Phe 250 | Gly | Asn | Leu | Arg | Ser 255 | Thr |
| Phe | Met | Ala | Leu 260 | Met | Ile | Gly | Ser | Tyr 265 | Ala | Ser | Ser | Ala | Ile 270 | Thr | Phe |
| Pro | Gly | Ile 275 | Lys | Leu | Ile | Tyr | Asp 280 | Ala | Gly | Val | Pro | Phe 285 | Thr | Val | Ile |
| Met | Phe 290 | Thr | Trp | Ser | Gly | Leu 295 | Ala | Cys | Leu | Ile | Phe 300 | Leu | Asn | Cys | Ala |
| Leu 305 | Asn | Trp | Pro | Ala | Glu 310 | Ala | Phe | Pro | Ala | Pro 315 | Glu | Glu | Val | Asp | Tyr 320 |
| Thr | Lys | Lys | Ile | Lys 325 | Leu | Ile | Gly | Leu | Ala 330 | Leu | Asp | His | Lys | Val 335 | Thr |

Gly Asp Arg Phe Tyr Thr His Val Thr Ile Val Gly Gln Arg Leu Ser Gln Lys Ser Pro Ser Leu Glu Glu Gly Ala Asp Ala Phe Ile Ser Ser Pro Asp Ile Pro Gly Thr Ser Glu Glu Thr Pro Glu Lys Ser Val Pro Phe Arg Lys Ser Leu Cys Ser Pro Ile Phe Leu Trp Ser Leu Val Thr Met Gly Met Thr Gln Leu Arg Val Ile Phe Tyr Met Gly Ala Met Asn Lys Ile Leu Glu Phe Ile Val Thr Gly Gly Lys Glu Arg Glu Thr Asn Glu Gln Arg Gln Lys Val Glu Glu Thr Val Glu Phe Tyr Ser Ser Ile Phe Gly Val Met Gln Leu Leu Cys Leu Leu Thr Cys Pro Leu Ile Gly Tyr Ile Met Asp Trp Arg Ile Lys Asp Cys Val Asp Ala Pro Thr Glu Gly Thr Leu Asn Glu Asn Ala Ser Phe Gly Asp Ala Arg Asp Gly Ala Ser Thr Lys Phe Thr Arg Pro Arg Tyr Arg Lys Val Gln Lys Leu Thr Asn Ala Ile Asn Ala Phe Thr Leu Thr Asn Ile Leu Leu Val Gly Phe Gly Ile Ala Cys Leu Ile Lys Asn Leu His Leu Gln Leu Leu Ala Phe Val Leu His Thr Ile Val Arg Gly Phe Phe His Ser Ala Cys Gly Gly Leu Tyr Ala Ala Val Phe Pro Ser Asn His Phe Gly Thr Leu Thr Gly Leu Gln Ser Leu Ile Ser Ala Val Phe Ala Leu Leu Gln Gln Leu Leu

Phe Met Ala Met Val Gly Pro Leu His Gly Asp Pro Phe Trp Val Asn 595 600 605

Leu Gly Leu Leu Leu Ser Phe Leu Gly Phe Leu Leu Pro Ser Tyr 610 620

Leu Tyr Tyr Tyr Arg Ser Arg Leu Gln Arg Glu Tyr Ala Thr Asn Leu 625 630 635 640

Val Asp Pro Gln Lys Val Leu Asn Thr Ser Lys Val Ala Thr 645 650

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<211> 401

<212> PRT

<213> Homo sapiens

<400> 367

Met Phe Gly Asn Leu Arg Ser Thr Leu Met Ala Leu Met Ile Gly Ser 1 5 10 15

Tyr Ala Ser Ser Ala Ile Thr Phe Pro Gly Ile Lys Leu Ile Tyr Asp 20 25 30

Ala Gly Val Ala Phe Val Val Ile Met Phe Thr Trp Ser Gly Leu Ala 35 40 45

Cys Leu Ile Phe Leu Asn Cys Thr Leu Asn Trp Pro Ile Glu Ala Phe 50 55 60

Pro Ala Pro Glu Glu Val Asn Tyr Thr Lys Lys Ile Lys Leu Ser Gly 65 70 75 80

Leu Ala Leu Asp His Lys Val Thr Gly Asp Leu Phe Tyr Thr His Val 85 90 95

Thr Thr Met Gly Gln Arg Leu Ser Gln Lys Ala Pro Ser Leu Glu Asp 100 105 110

Gly Ser Asp Ala Phe Met Ser Pro Gln Asp Val Arg Gly Thr Ser Glu 115 120 125

Asn Leu Pro Glu Arg Ser Val Pro Leu Arg Lys Ser Leu Cys Ser Pro 130 135 140

Thr Phe Leu Trp Ser Leu Leu Thr Met Gly Met Thr Gln Leu Arg Ile

- Ile Phe Tyr Met Ala Ala Val Asn Lys Met Leu Glu Tyr Leu Val Thr 165 170 175
- Gly Gly Gln Glu His Glu Thr Asn Glu Gln Gln Gln Lys Val Ala Glu 180 185 190
- Thr Val Gly Phe Tyr Ser Ser Val Phe Gly Ala Met Gln Leu Leu Cys 195 200 205
- Leu Leu Thr Cys Pro Leu Ile Gly Tyr Ile Met Asp Trp Arg Ile Lys 210 215 220
- Asp Cys Val Asp Ala Pro Thr Gln Gly Thr Val Leu Gly Asp Ala Arg 225 230 235 240
- Asp Gly Val Ala Thr Lys Ser Ile Arg Pro Arg Tyr Cys Lys Ile Gln 245 250 255
- Lys Leu Thr Asn Ala Ile Ser Ala Phe Thr Leu Thr Asn Leu Leu Leu 260 265 270
- Val Gly Phe Gly Ile Thr Cys Leu Ile Asn Asn Leu His Leu Gln Phe 275 280 285
- Val Thr Phe Val Leu His Thr Ile Val Arg Gly Phe Phe His Ser Ala 290 295 300
- Cys Gly Ser Leu Tyr Ala Ala Val Phe Pro Ser Asn His Phe Gly Thr 305 310 315
- Leu Thr Gly Leu Gln Ser Leu Ile Ser Ala Val Phe Ala Leu Leu Gln 325 330 335
- Gln Pro Leu Phe Met Ala Met Val Gly Pro Leu Lys Gly Glu Pro Phe 340 345 350
- Trp Val Asn Leu Gly Leu Leu Leu Phe Ser Leu Leu Gly Phe Leu Leu 355 360 365
- Pro Ser Tyr Leu Phe Tyr Tyr Arg Ala Arg Leu Gln Gln Glu Tyr Ala 370 375 380
- Ala Asn Gly Met Gly Pro Leu Lys Val Leu Ser Gly Ser Glu Val Thr 385 390 395 400

Ala

<210> 368 <211> 489 <212> PRT <213> Homo sapiens <400> 368 Met Ala Pro Thr Leu Ala Thr Ala His Arg Arg Trp Trp Met Ala Cys Thr Pro Val Leu Glu Asn Leu Leu Phe Ser Ala Val Leu Leu Gly Trp Gly Ser Leu Leu Ile Met Leu Lys Ser Glu Gly Phe Tyr Ser Tyr Leu Cys Thr Glu Pro Glu Asn Val Thr Asn Gly Thr Val Gly Gly Thr Ala Glu Pro Gly His Glu Glu Val Ser Trp Met Asn Gly Trp Leu Ser Cys Gln Ala Gln Asp Glu Met Leu Asn Leu Ala Phe Thr Val Gly Ser Phe Leu Leu Ser Ala Ile Thr Leu Pro Leu Gly Ile Val Met Asp Lys Tyr Gly Pro Arg Lys Leu Arg Leu Leu Gly Ser Ala Cys Phe Ala Val Ser Cys Leu Leu Ile Ala Tyr Gly Ala Ser Lys Pro Asn Ala Leu Ser Val Leu Ile Phe Ile Ala Leu Ala Leu Asn Gly Phe Gly Met Cys Met Thr Phe Thr Ser Leu Thr Leu Pro Asn Met Phe Gly Asp Leu Arg Ser Thr Phe Ile Ala Leu Met Ile Gly Ser Tyr Ala Ser Ser Ala Val

Thr Phe Pro Gly Ile Lys Leu Ile Tyr Asp Ala Gly Val Ser Phe Ile

| Val | Val 210 | Leu | Val | Val | Trp | Ala 215 | Gly | Cys | Ser | Gly | Leu 220 | Val | Phe | Leu | Asn |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Cys 225 | Phe | Phe | Asn | Trp | Pro 230 | Leu | Glu | Pro | Phe | Pro 235 | Gly | Pro | Glu | Asp | Met 240 |
| Asp | Tyr | Ser | Val | Lys 245 | Ile | Lys | Phe | Ser | Trp 250 | Leu | Gly | Phe | Asp | His 255 | Lys |
| Ile | Thr | Gly | Lys 260 | Gln | Phe | Tyr | Lys | Gln 265 | Val | Thr | Thr | Val | Gly 270 | Arg | Arg |
| Leu | Ser | Val 275 | Gly | Ser | Ser | Met | Arg 280 | Ser | Ala | Lys | Glu | Gln 285 | Val | Ala | Leu |
| Gln | Glu 290 | Gly | His | Lys | Leu | Cys 295 | Leu | Ser | Thr | Val | Asp 300 | Leu | Glu | Val | Lys |
| Cys 305 | Gln | Pro | Asp | Ala | Ala 310 | Val | Ala | Pro | Ser | Phe 315 | Met | His | Ser | Val | Phe 320 |
| Ser | Pro | Ile | Leu | Leu 325 | Leu | Ser | Leu | Val | Thr 330 | Met | Cys | Val | Thr | Gln 335 | Leu |
| Arg | Leu | Ile | Phe 340 | Tyr | Met | Gly | Ala | Met 345 | Asn | Asn | Ile | Leu | Lys 350 | Phe | Leu |
| Val | Ser | Gly 355 | Asp | Gln | Lys | Thr | Val 360 | Gly | Leu | Tyr | Thr | Ser 365 | Ile | Phe | Gly |
| Val | Leu 370 | Gln | Leu | Leu | Cys | Leu 375 | Leu | Thr | Ala | Pro | Val 380 | Ile | Gly | Tyr | Ile |
| Met 385 | Asp | Trp | Arg | Leu | Lys 390 | Glu | Cys | Glu | Asp | Ala 395 | Ser | Glu | Glu | Pro | Glu 400 |
| Glu | Lys | Asp | Ala | Asn 405 | Gln | Cys | Val | Gly | Arg 410 | Ala | Gly | Ala | Pro | Ala 415 | Pro |
| Ser | Pro | Gln | Pro 420 | Leu | Gln | Lys | Asp | Pro 425 | Arg | Ala | Ala | Cys | Gln 430 | Ala | Gln |
| Gly | Gly | Trp 435 | Asp | Arg | Gly | Arg | Glu 440 | Gln | Cys | Thr | Pro | Ala 445 | Pro | Pro | Gly |
| Ala | Leu 450 | Arg | Glu | Ala | His | Ser 455 | Phe | Ser | Ser | Ala | Cys 460 | Val | Ser | Thr | Ala |

Pro Leu Phe Met Glu Ile Val Trp Asn Ala Met Glu Met Leu Glu Phe 465 470 475 480

Glu Ala Arg Cys Gly Asp Ser Cys Leu 485

<210> 369

<211> 373

<212> PRT

<213> Homo sapiens

<400> 369

Ile Lys Leu Ile Tyr Asp Ala Gly Val Ser Phe Ile Val Val Leu Val 1 5 10 15

Val Trp Ala Gly Cys Ser Gly Leu Val Phe Leu Asn Cys Phe Phe Asn 20 25 30

Trp Pro Leu Glu Pro Phe Pro Gly Pro Glu Asp Met Asp Tyr Ser Val
35 40 45

Lys Ile Lys Phe Ser Trp Leu Gly Phe Asp His Lys Ile Thr Gly Lys 50 55 60

Gln Phe Tyr Lys Gln Val Thr Thr Val Gly Arg Arg Leu Ser Val Gly 65 70 75 80

Ser Ser Met Arg Ser Ala Lys Glu Gln Val Ala Leu Gln Glu Gly His
85 90 95

Lys Leu Cys Leu Ser Thr Val Asp Leu Glu Val Lys Cys Gln Pro Asp 100 105 110

Ala Ala Val Val Pro Ser Phe Met His Ser Val Phe Ser Pro Ile Leu 115 120 125

Leu Leu Ser Leu Val Thr Met Cys Val Thr Gln Leu Arg Leu Ile Phe 130 135 140

Gln Lys Thr Val Gly Leu Tyr Thr Ser Ile Phe Gly Val Leu Gln Leu 165 170 175

Leu Cys Leu Leu Thr Ala Pro Val Ile Gly Tyr Ile Met Asp Trp Arg 180 185 190

Leu Lys Glu Cys Glu Asp Ala Ser Glu Glu Pro Glu Glu Lys Asp Ala 195 200 Asn Gln Gly Glu Lys Lys Lys Lys Arg Asp Arg Gln Ile Gln Lys 210 215 220 Ile Thr Asn Ala Met Arg Ala Phe Ala Phe Thr Asn Leu Leu Val 225 230 235 Gly Phe Gly Val Thr Cys Leu Ile Pro Asn Leu Pro Leu Gln Ile Leu 250 245 Ser Phe Ile Leu His Thr Ile Val Arg Gly Phe Ile His Ser Ala Val 265 260 Gly Gly Leu Tyr Ala Ala Val Tyr Pro Ser Thr Gln Phe Gly Ser Leu 280 Thr Gly Leu Gln Ser Leu Ile Ser Ala Leu Phe Ala Leu Leu Gln Gln 295 300 Pro Leu Phe Leu Ala Met Met Gly Pro Leu Gln Gly Asp Pro Leu Trp 310 315 Val Asn Val Gly Leu Leu Leu Ser Leu Leu Gly Phe Cys Leu Pro 325 330 335 Leu Tyr Leu Ile Cys Tyr Arg Arg Gln Leu Glu Arg Gln Leu Gln Gln 345 340 350 Arg Gln Glu Asp Asp Lys Leu Phe Leu Lys Ile Asn Gly Ser Ser Asn 355 360 365 Gln Glu Ala Phe Val 370 <210> 370 <211> 125 <212> PRT <213> Homo sapiens

<400> 370

Met Ala Gly Pro Ser Leu Ala Cys Cys Leu Leu Gly Leu Leu Ala Leu
1 5 10 15

Thr Ser Ala Cys Tyr Ile Gln Asn Cys Pro Leu Gly Gly Lys Arg Ala

| 20 | 25 | 30 |
|----|----|----|
| | | |

Ala Pro Asp Leu Asp Val Arg Lys Cys Leu Pro Cys Gly Pro Gly Gly
35 40 45

Lys Gly Arg Cys Phe Gly Pro Asn Ile Cys Cys Ala Glu Glu Leu Gly 50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Ala Cys Gly Ser Gly Gly 85 90 95

Arg Cys Ala Val Leu Gly Leu Cys Cys Ser Pro Asp Gly Cys His Ala 100 105 110

Asp Pro Ala Cys Asp Ala Glu Ala Thr Phe Ser Gln Arg 115 120 125

<210> 371

<211> 124

<212> PRT

<213> Homo sapiens

<400> 371

Met Ala Gly Pro Ser Leu Ala Cys Cys Leu Leu Gly Leu Leu Ala Leu 1 5 10 15

Thr Ser Ala Cys Tyr Ile Gln Asn Cys Pro Leu Gly Gly Lys Arg Ala 20 25 30

Ala Pro Asp Leu Asp Val Arg Lys Cys Leu Pro Cys Gly Pro Gly Gly
35 40 45

Lys Gly Arg Cys Phe Gly Pro Asn Ile Cys Cys Ala Glu Glu Leu Gly
50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Ala Cys Gly Ser Gly Gly 85 90 95

Arg Cys Ala Leu Gly Leu Cys Cys Ser Pro Asp Gly Cys His Ala Asp 100 105 110 Pro Ala Cys Asp Ala Glu Ala Thr Phe Ser Gln Arg 115 120

<210> 372

<211> 125

<212> PRT

<213> Sus scrofa

<400> 372

O

Met Ala Gly Pro Ser Leu Ala Cys Cys Leu Leu Gly Leu Leu Ala Leu 1 5 10 15

Thr Ser Ala Cys Tyr Ile Gln Asn Cys Pro Leu Gly Gly Lys Arg Ala
20 25 30

Val Leu Asp Leu Asp Val Arg Lys Cys Leu Pro Cys Gly Pro Gly Gly
35 40 45

Lys Gly Arg Cys Phe Gly Pro Ser Ile Cys Cys Gly Asp Glu Leu Gly 50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Pro Cys Gly Ser Glu Gly
85 90 95

Arg Cys Ala Ala Ala Gly Ile Cys Cys Asn Pro Asp Gly Cys Arg Phe 100 105 110

Asp Pro Ala Cys Asp Pro Glu Ala Thr Phe Ser Gln Arg 115 120 125

<210> 373

<211> 125

<212> PRT

<213> Ovis aries

<400> 373

Met Ala Gly Ser Ser Leu Ala Cys Cys Leu Leu Gly Leu Leu Ala Leu 1 5 10 15

Thr Ser Ala Cys Tyr Ile Gln Asn Cys Pro Leu Gly Gly Lys Arg Ala
20 25 30

Val Leu Asp Leu Asp Val Arg Thr Cys Leu Pro Cys Gly Pro Gly Gly

35 40 45

Lys Gly Arg Cys Phe Gly Pro Ser Ile Cys Cys Gly Asp Glu Leu Gly 50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Arg Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Pro Cys Gly Ser Gly Gly
85 90 95

Arg Cys Ala Ala Ala Gly Ile Cys Cys Ser Pro Asp Gly Cys His Ala 100 105 110

Asp Pro Ala Cys Asp Pro Glu Ala Ala Phe Ser Gln His
115 120 125

<210> 374

<211> 125

<212> PRT

<213> Bos taurus

<400> 374

Met Ala Gly Ser Ser Leu Ala Cys Cys Leu Leu Gly Leu Leu Ala Leu 1 5 10 15

Thr Ser Ala Cys Tyr Ile Gln Asn Cys Pro Leu Gly Gly Lys Arg Ala
20 25 30

Val Leu Asp Leu Asp Val Arg Thr Cys Leu Pro Cys Gly Pro Gly Gly
35 40 45

Lys Gly Arg Cys Phe Gly Pro Ser Ile Cys Cys Gly Asp Glu Leu Gly 50 55 60

Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln Glu Glu Asn Tyr 65 70 75 80

Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Pro Cys Gly Ser Gly Gly
85 90 95

Arg Cys Ala Ala Gly Ile Cys Cys Ser Pro Asp Gly Cys His Glu
100 105 110

Asp Pro Ala Cys Asp Pro Glu Ala Ala Phe Ser Gln His
115 120 125

<210> 375

<211> 56

<212> PRT

<213> Homo sapiens

<400> 375

Glu Glu Leu Gly Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln
1 5 10 15

Glu Glu Asn Tyr Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Ala Cys
20 25 30

Gly Ser Gly Gly Arg Cys Ala Val Leu Gly Leu Cys Cys Ser Pro Asp 35 40 45

Gly Cys His Ala Asp Pro Ala Cys
50 55

<210> 376

<211> 57

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence:
Neurohypophysial hormones domain sequence

<400> 376

Glu Glu Leu Gly Cys Tyr Val Gly Thr Pro Glu Thr Ala Arg Cys Gln
1 5 10 15

Glu Glu Asn Tyr Leu Pro Ser Pro Cys Glu Ala Gly Gly Lys Pro Cys
20 25 30

Gly Ser Asp Ala Gly Arg Cys Ala Ala Pro Gly Val Cys Cys Asp Ser 35 40 45

Glu Ser Cys Val Val Asp Pro Glu Cys
50 55

<210> 377

<211> 56

<212> PRT

<213> Homo sapiens

<400> 377 Glu Glu Leu Gly Cys Phe Val Gly Thr Ala Glu Ala Leu Arg Cys Gln 10 Glu Glu Asn Tyr Leu Pro Ser Pro Cys Gln Ser Gly Gln Lys Ala Cys 20 25 Gly Ser Gly Gly Arg Cys Ala Val Leu Gly Leu Cys Cys Ser Pro Asp 40 45 Gly Cys His Ala Asp Pro Ala Cys 50 55 <210> 378 <211> 57 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Neurohypophysial hormones domain sequence <400> 378 Glu Glu Leu Gly Cys Tyr Val Gly Thr Pro Glu Thr Ala Arg Cys Gln 10 Glu Glu Asn Tyr Leu Pro Ser Pro Cys Glu Ser Gly Gly Arg Pro Cys 20 25 Gly Ser Asp Gly Gly Arg Cys Ala Ala Pro Gly Ile Cys Cys Asp Ser Glu Ser Cys Ala Ala Asp Pro Ser Cys 50 55 <210> 379 <211> 158 <212> PRT <213> Homo sapiens <400> 379 Met Ser Asp Lys Ser Asn Met Asp Glu Ile Glu Lys Phe Ser Lys Ser 1 5 10 15

20

Lys Leu Lys Lys Thr Glu Met Gln Glu Lys Asn Pro Gln Pro Ser Lys

25

Glu Trp Ile Glu Gln Glu Lys Gln Ala Gly Phe Cys Ala Met Ala Ala 35 40 45

Asn Ser Ser Phe Leu Gly Gly Val His Gly Leu Phe Leu Val Trp Val 50 55 60

Ala Leu Arg Val Leu Gly Asp Arg Pro Phe Lys Cys Thr Phe Met Ser 65 70 75 80

Leu Thr Leu His Tyr Pro Arg Cys Arg Leu Glu Thr Gly Ile Gln Gly
85 90 95

Ala Phe Gly Lys Pro Gln Gly Thr Val Ala Arg Val His Ile Gly Gln
100 105 110

Val Lys Ser Ile Cys Thr Lys Leu Gln Asn Lys Glu His Val Ile Glu 115 120 125

Ala Pro Cys Arg Ala Lys Phe Lys Phe Pro Gly His Gln Lys Ile His 130 135 140

Ile Ser Lys Lys Trp Gly Phe Thr Lys Phe Asn Val Asp Glu 145 150 155

<210> 380

<211> 56

<212> PRT

<213> Rattus norvegicus

<400> 380

Leu Phe Ala Gln Leu Ala Gln Leu Leu Pro Ala Thr Met Ser Asp Lys
1 5 10 15

Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser Lys Leu Lys Lys
20 25 30

Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu Thr Ile Glu
35 40 45

Gln Glu Lys Gln Ala Gly Glu Ser 50 55

<210> 381

<211> 50

<212> PRT

<210> 382 <211> 43 <212> PRT <213> Orycctolagus cuniculus

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys Glu
20 25 30

Thr Ile Glu Gln Glu Lys Gln Ala Gly Glu Ser 35 40

<210> 383 <211> 44 <212> PRT <213> Mus musculus

<400> 383

Met Ser Asp Lys Pro Asp Met Ala Glu Ile Glu Lys Phe Asp Lys Ser 1 5 10 15

Lys Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Ser Lys
20 25 30

Glu Thr Ile Glu Gln Glu Lys Gln Ala Gly Glu Ser 35 40

```
<400> 384
Met Asp Glu Ile Glu Lys Phe Ser Lys Ser Lys Leu Lys Lys Thr Glu
 1
                  5
                                      10
                                                          15
Met Gln Glu Lys Asn Pro Gln Pro Ser Lys Glu Trp Ile Glu Gln Glu
                                  25
Lys Gln Ala Gly
         35
<210> 385
<211> 36
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Thymosin beta
      actin-binding motif sequence
<400> 385
Thr Asp Glu Ile Glu Asn Phe Asp Ser Glu Asn Leu Lys Lys Thr Glu
                  5
                                      10
                                                          15
Thr Ile Glu Lys Asn Val Leu Pro Ser Lys Glu Asp Ile Glu Gln Glu
                                  25
             20
                                                      30
Lys Gln Leu Gln
         35
<210> 386
<211> 41
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Thymosin
      beta-4 family domain sequence
<400> 386
Ser Asp Lys Pro Asp Leu Glu Glu Ile Ala Ser Phe Asp Lys Ala Lys
                                      10
                                                           15
```

<210> 384 <211> 36 <212> PRT

<213> Homo sapiens

Leu Lys Lys Thr Glu Thr Gln Glu Lys Asn Pro Leu Pro Thr Lys Glu
20 25 30

Thr Ile Glu Gln Glu Lys Gln Ala Glu 35 40

<210> 387

<211> 36

<212> PRT

<213> Homo sapiens

<400> 387

Met Asp Glu Ile Glu Lys Phe Ser Lys Ser Lys Leu Lys Lys Thr Glu
1 5 10 15

Met Gln Glu Lys Asn Pro Gln Pro Ser Lys Glu Trp Ile Glu Gln Glu 20 25 30

Lys Gln Ala Gly 35

<210> 388

<211> 132

<212> PRT

<213> Mus musculus

<400> 388

Met Val Asp Gln Leu Gln Gly Thr Trp Lys Ser Val Ser Cys Asp Asn 1 5 10 15

Phe Glu Asn Tyr Met Lys Glu Leu Gly Val Gly Arg Ala Ser Arg Lys 20 25 30

Leu Gly Cys Leu Ala Lys Pro Thr Val Thr Ile Ser Thr Asp Gly Asp 35 40 45

Leu Ile Thr Ile Lys Thr Lys Ser Ile Phe Lys Asn Lys Glu Ile Ser 50 55 60

Phe Lys Leu Gly Glu Glu Phe Glu Glu Thr Thr Pro Ser Gly Arg Lys
65 70 75 80

Ser Lys Ser Thr Val Ile Leu Asp Asn Asp Ser Leu Val Gln Val Gln 85 90 95

Asp Trp Asp Gly Lys Glu Ala Thr Ile Cys Arg Arg Leu Val Asp Gly 100 105 110

Lys Met Val Val Glu Ser Ala Val Asn Asn Val Thr Cys Thr Arg Thr 115 120 125

Tyr Gln Arg Val 130

<210> 389

<211> 132

<212> PRT

<213> Orycctolagus cuniculus

<400> 389

Met Ser Asn Lys Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 1 5 10 15

Phe Asp Asp Tyr Met Lys Ala Leu Gly Val Gly Leu Ala Thr Arg Lys
20 25 30

Leu Gly Asn Leu Ala Lys Pro Asn Val Ile Ile Ser Lys Lys Gly Asp 35 40 45

Ile Ile Thr Ile Arg Thr Glu Ser Thr Phe Lys Asn Thr Glu Ile Ser
50 55 60

Phe Lys Leu Gly Gln Glu Phe Glu Glu Thr Thr Ala Asp Asn Arg Lys 65 70 75 80

Thr Lys Ser Ile Ile Thr Leu Glu Arg Gly Ala Leu Asn Gln Val Gln 85 90 95

Lys Trp Asp Gly Lys Glu Thr Thr Ile Lys Arg Lys Leu Val Asp Gly 100 105 110

Lys Met Val Val Glu Cys Lys Met Lys Gly Val Val Cys Thr Arg Ile 115 120 125

Tyr Glu Lys Val 130

<210> 390

<211> 132

<212> PRT

<213> Homo sapiens

<400> 390

Met Ser Asn Lys Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn
1 5 10 15

Phe Asp Asp Tyr Met Lys Ala Leu Gly Val Gly Leu Ala Thr Arg Lys
20 25 30

Leu Gly Asn Leu Ala Lys Pro Thr Val Ile Ile Ser Lys Lys Gly Asp 35 40 45

Ile Ile Thr Ile Arg Thr Glu Ser Thr Phe Lys Asn Thr Glu Ile Ser 50 55 60

Phe Lys Leu Gly Gln Glu Phe Glu Glu Thr Thr Ala Asp Asn Arg Lys
65 70 75 80

Thr Lys Ser Ile Val Thr Leu Gln Arg Gly Ser Leu Asn Gln Val Gln 85 90 95

Arg Trp Asp Gly Lys Glu Thr Thr Ile Lys Arg Lys Leu Val Asn Gly
100 105 110

Lys Met Val Ala Glu Cys Lys Met Lys Gly Val Val Cys Thr Arg Ile 115 120 125

Tyr Glu Lys Val 130

<210> 391

<211> 132

<212> PRT

<213> Mus musculus

<400> 391

Met Ser Asn Lys Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu His

1 5 10 15

Phe Asp Asp Tyr Met Lys Ala Leu Gly Val Gly Leu Ala Asn Arg Lys 20 25 30

Leu Gly Asn Leu Ala Lys Pro Thr Val Ile Ile Ser Lys Lys Gly Asp 35 40 45

Tyr Ile Thr Ile Arg Thr Glu Ser Ala Phe Lys Asn Thr Glu Ile Ser 50 55 60

Phe Lys Leu Gly Gln Glu Phe Asp Glu Thr Thr Ala Asp Asn Arg Lys
65 70 75 80

Ala Lys Ser Ile Val Thr Leu Glu Arg Gly Ser Leu Lys Gln Val Gln 85 90 95

Lys Trp Asp Gly Lys Glu Thr Ala Ile Arg Arg Thr Leu Leu Asp Gly
100 105 110

Arg Met Val Val Glu Cys Ile Met Lys Gly Val Val Cys Thr Arg Ile 115 120 125

Tyr Glu Lys Val 130

<210> 392

<211> 132

<212> PRT

<213> Bos taurus

<400> 392

Met Cys Asp Ala Phe Val Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 1 5 10 15

Phe Asp Asp Tyr Met Lys Glu Val Gly Val Gly Phe Ala Thr Arg Lys
20 25 30

Val Ala Gly Met Ala Lys Pro Thr Leu Ile Ile Ser Leu Asn Gly Gly
35 40 45

Val Val Thr Ile Lys Ser Glu Ser Thr Phe Lys Asn Thr Glu Ile Ser 50 55 60

Phe Lys Leu Gly Gln Glu Phe Asp Glu Ile Thr Pro Asp Asp Arg Lys
65 70 75 80

Val Lys Ser Ile Val Asn Leu Asp Glu Gly Ala Leu Val Gln Val Gln 85 90 95

Asn Trp Asp Gly Lys Ser Thr Thr Ile Lys Arg Lys Leu Met Asp Asp 100 105 110

Lys Met Val Leu Glu Cys Val Met Asn Gly Val Thr Ala Thr Arg Val 115 120 125

Tyr Glu Arg Ala

130

<210> 393 <211> 129 <212> PRT <213> Homo sapiens <4.00> 393 Gln Leu Gln Gly Thr Trp Lys Ser Ile Ser Cys Glu Asn Ser Glu Asp 5 10 Tyr Met Lys Glu Leu Gly Ile Gly Arg Ala Ser Arg Lys Leu Gly Arg 20 25 30 Leu Ala Lys Pro Thr Val Thr Ile Ser Thr Asp Gly Asp Val Ile Thr 35 40 45 Ile Lys Thr Lys Ser Ile Phe Lys Asn Asn Glu Ile Ser Phe Lys Leu 50 55 60 Gly Glu Glu Phe Glu Glu Ile Thr Pro Gly Gly His Lys Thr Lys Ser 65 70 75 80 Lys Val Thr Leu Asp Lys Glu Ser Leu Ile Gln Val Gln Asp Trp Asp 85 90 95 Gly Lys Glu Thr Thr Ile Thr Arg Lys Leu Val Asp Gly Lys Met Val 100 105 110 Val Glu Ser Thr Val Asn Ser Val Ile Cys Thr Arg Thr Tyr Glu Lys 115 120 125 Val <210> 394 <211> 145 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: lipocalin domain sequence <400> 394 Lys Phe Ala Gly Lys Trp Tyr Leu Val Ala Ser Ala Asn Phe Asp Pro

10

15

5

1

Glu Leu Lys Glu Glu Leu Gly Val Leu Glu Ala Thr Arg Lys Glu Ile 30 20 25 Thr Pro Leu Lys Glu Gly Asn Leu Glu Ile Val Phe Asp Gly Asp Lys 40 35 Asn Gly Ile Cys Glu Glu Thr Phe Gly Lys Leu Glu Lys Thr Lys Lys 55 Leu Gly Val Glu Phe Asp Tyr Tyr Thr Gly Asp Asn Arg Phe Val Val 70 Leu Asp Thr Asp Tyr Asp Asn Tyr Leu Leu Val Cys Val Gln Lys Gly 85 90 Asp Gly Asn Glu Thr Ser Arg Thr Ala Glu Leu Tyr Gly Arg Thr Pro 105 Glu Leu Ser Pro Glu Ala Leu Glu Leu Phe Glu Thr Ala Thr Lys Glu 120 Leu Gly Ile Pro Glu Asp Asn Val Val Cys Thr Arg Gln Thr Glu Arg 135 140 Cys 145 <210> 395 <211> 132 <212> PRT <213> Homo sapiens <400> 395 Met Val Glu Pro Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 10 Phe Glu Asp Tyr Met Lys Glu Leu Gly Val Asn Phe Ala Ala Arg Asn 30 Met Ala Gly Leu Val Lys Pro Thr Val Thr Ile Ser Val Asp Gly Lys 35 40

55

Met Met Thr Ile Arg Thr Glu Ser Ser Phe Gln Asp Thr Lys Ile Ser

Phe Lys Leu Gly Glu Glu Phe Asp Glu Thr Thr Ala Asp Asn Arg Lys

Val Lys Ser Thr Ile Thr Leu Glu Asn Gly Ser Met Ile His Val Gln 85 90 95

Lys Trp Leu Gly Lys Glu Thr Thr Ile Lys Arg Lys Ile Val Asp Glu 100 105 110

Lys Met Val Val Glu Cys Lys Met Asn Asn Ile Val Ser Thr Arg Ile 115 120 125

Tyr Glu Lys Val 130

<210> 396

<211> 132

<212> PRT

<213> Mus musculus

<400> 396

Met Ile Glu Pro Phe Leu Gly Thr Trp Lys Leu Ile Ser Ser Glu Asn
1 5 10 15

Phe Glu Asn Tyr Val Arg Glu Leu Gly Val Glu Cys Glu Pro Arg Lys
20 25 30

Val Ala Cys Leu Ile Lys Pro Ser Val Ser Ile Ser Phe Asn Gly Glu 35 40 45

Arg Met Asp Ile Gln Ala Gly Ser Ala Cys Arg Asn Thr Glu Ile Ser 50 55 60

Phe Lys Leu Gly Glu Glu Phe Glu Glu Thr Thr Ala Asp Asn Arg Lys 65 70 75 80

Val Lys Ser Leu Ile Thr Phe Glu Gly Gly Ser Met Ile Gln Val Gln 85 90 95

Lys Trp Leu Gly Lys Gln Thr Thr Ile Lys Arg Lys Ile Val Asp Gly
100 105 110

Lys Met Val Val Glu Cys Thr Met Asn Asn Val Val Ser Thr Arg Ile 115 120 125

Tyr Glu Arg Val

<210> 397 <211> 132 <212> PRT <213> Mus musculus <400> 397 Met Ile Glu Pro Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 10 Phe Glu Asn Tyr Val Arg Glu Leu Gly Val Glu Cys Glu Pro Arg Lys 20 25 Val Ala Cys Leu Ile Lys Pro Ser Val Ser Ile Ser Phe Asn Gly Glu 40 Arg Met Asp Ile Gln Ala Gly Ser Ala Cys Arg Asn Thr Lys Ile Ser Phe Lys Leu Gly Glu Glu Phe Glu Glu Thr Thr Ala Asp Asn Arg Lys 70 75 Val Lys Ser Leu Ile Thr Phe Glu Gly Gly Ser Met Ile Gln Ile Gln 85 90 Arg Trp Leu Gly Lys Gln Thr Thr Ile Lys Arg Arg Ile Val Asp Gly 105 100 Arg Met Val Val Glu Cys Thr Met Asn Asn Val Val Ser Thr Arg Thr 115 120

125

110

Tyr Glu Arg Val 130

<210> 398

<211> 132

<212> PRT

<213> Rattus norvegicus

<400> 398

Met Ile Glu Pro Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn 10

Phe Glu Asn Tyr Val Arg Glu Leu Gly Val Glu Cys Glu Pro Arg Lys 25

Val Ala Cys Leu Ile Lys Pro Ser Val Ser Ile Ser Phe Asn Gly Glu

35 40 45

Arg Met Asp Ile Gln Ala Gly Ser Ala Cys Arg Asn Thr Glu Ile Ser 50 55 60

Phe Lys Leu Gly Glu Glu Phe Glu Glu Thr Thr Ala Asp Asn Arg Lys
65 70 75 80

Val Lys Ser Leu Ile Thr Phe Glu Gly Gly Ser Met Ile Gln Ile Gln 85 90 95

Arg Trp Leu Gly Lys Gln Thr Thr Ile Lys Arg Arg Ile Val Asp Gly \$100\$

Arg Met Val Val Glu Cys Thr Met Asn Asn Val Val Ser Thr Arg Thr 115 120 125

Tyr Glu Arg Val 130

<210> 399

<211> 132

<212> PRT

<213> Sus scrofa

<400> 399

Met Cys Asp Ala Phe Val Gly Thr Trp Lys Leu Val Ser Ser Glu Asn $1 \hspace{1cm} 5 \hspace{1cm} 10 \hspace{1cm} 15$

Phe Asp Asp Tyr Met Lys Glu Val Gly Val Gly Phe Ala Thr Arg Lys
20 25 30

Val Ala Gly Met Ala Lys Pro Asn Leu Ile Ile Thr Val Asn Gly Asp 35 40 45

Met Ile Thr Ile Arg Ser Glu Ser Thr Phe Lys Asn Thr Glu Ile Ala 50 55 60

Phe Lys Leu Gly Gln Glu Phe Asp Glu Val Thr Ala Asp Asp Arg Lys 65 70 75 80

Val Lys Ser Thr Ile Thr Leu Asp Gly Gly Ala Leu Val Gln Val Gln 85 90 95

Lys Trp Asp Gly Lys Thr Thr Thr Ile Asn Arg Lys Ile Val Asp Asp 100 105 110

Lys Leu Val Val Glu Cys Ile Met Lys Gly Val Thr Ala Thr Arg Ile 115 120 125

Tyr Glu Arg Ala 130

<210> 400

<211> 124

<212> PRT

<213> Homo sapiens

<400> 400

Phe Leu Gly Thr Trp Lys Leu Val Ser Ser Glu Asn Phe Glu Asp Tyr

1 5 10 15

Met Lys Glu Leu Gly Phe Ala Ala Arg Asn Met Ala Gly Leu Val Lys
20 25 30

Pro Thr Val Thr Ile Ser Val Asp Gly Lys Met Met Thr Ile Arg Thr 35 40 45

Glu Ser Ser Phe Gln Asp Thr Lys Ile Ser Phe Lys Leu Gly Glu Glu 50 55 60

Phe Asp Glu Thr Thr Ala Asp Asn Arg Lys Val Lys Ser Thr Ile Thr 65 70 75 80

Leu Glu Asn Gly Ser Met Ile His Val Gln Lys Trp Leu Gly Lys Glu 85 90 95

Thr Thr Ile Lys Arg Lys Ile Val Asp Glu Lys Met Val Val Glu Cys 100 105 110

Lys Met Asn Asn Ile Val Ser Thr Arg Ile Tyr Glu 115 120

<210> 401

<211> 127

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: lipocalin domain sequence

<400> 401

Phe Ala Gly Lys Trp Tyr Leu Val Ala Ser Ala Asn Phe Asp Pro Glu Leu Lys Glu Glu Leu Gly Val Leu Glu Ala Thr Arg Lys Glu Ile Thr Pro Leu Lys Glu Gly Asn Leu Glu Ile Val Phe Asp Gly Asp Lys Asn Gly Ile Cys Glu Glu Thr Phe Gly Lys Leu Glu Lys Thr Lys Leu Gly Val Glu Phe Asp Tyr Tyr Thr Gly Asp Asn Arg Phe Val Val Leu Asp Thr Asp Tyr Asp Asn Tyr Leu Leu Val Cys Val Gln Lys Gly Asp Gly Asn Glu Thr Ser Arg Thr Ala Glu Leu Tyr Gly Arg Thr Pro Glu Leu Ser Pro Glu Ala Leu Glu Leu Phe Glu Thr Ala Thr Lys Glu <210> 402 <211> 391 <212> PRT <213> Homo sapiens <400> 402 His Gln Ala Ala His Gln Pro Phe Pro Arg Pro Arg Phe Arg Gln Glu Thr Gly His Pro Ser Leu Gln Arg Asp Phe Pro Arg Ser Phe Leu Leu Asp Leu Pro Asn Phe Pro Asp Leu Ser Lys Ala Asp Ile Asn Gly Gln Asn Pro Asn Ile Gln Val Thr Ile Glu Val Val Asp Gly Pro Asp Ser Glu Ala Asp Lys Asp Gln His Pro Glu Asn Lys Pro Ser Trp Ser Val Pro Ser Pro Asp Trp Arg Ala Trp Trp Gln Arg Ser Leu Ser Leu Ala

Arg Ala Asn Ser Gly Asp Gln Asp Tyr Lys Tyr Asp Ser Thr Ser Asp Asp Ser Asn Phe Leu Asn Pro Pro Arg Gly Trp Asp His Thr Ala Pro Gly His Arg Thr Phe Glu Thr Lys Asp Gln Pro Glu Tyr Asp Ser Thr Asp Gly Glu Gly Asp Trp Ser Leu Trp Ser Val Cys Ser Val Thr Cys Gly Asn Gly Asn Gln Lys Arg Thr Arg Ser Cys Gly Tyr Ala Cys Thr Ala Thr Glu Ser Arg Thr Cys Asp Arg Pro Asn Cys Pro Gly Ile Glu Asp Thr Phe Arg Thr Ala Ala Thr Glu Val Ser Leu Leu Ala Gly Ser Glu Glu Phe Asn Ala Thr Lys Leu Phe Glu Val Asp Thr Asp Ser Cys Glu Arg Trp Met Ser Cys Lys Ser Glu Phe Leu Lys Lys Tyr Met His Lys Val Met Asn Asp Leu Pro Ser Cys Pro Cys Ser Tyr Pro Thr Glu Val Ala Tyr Ser Thr Ala Asp Ile Phe Asp Arg Ile Lys Arg Lys Asp Phe Arg Trp Lys Asp Ala Ser Gly Pro Lys Glu Lys Leu Glu Ile Tyr Lys Pro Thr Ala Arg Tyr Cys Ile Arg Ser Met Leu Ser Leu Glu Ser Thr Thr Leu Ala Ala Gln His Cys Cys Tyr Gly Asp Asn Met Gln Leu Ile Thr Arg Gly Lys Gly Ala Gly Thr Pro Asn Leu Ile Ser Thr Glu Phe Ser Ala Glu Leu His Tyr Lys Val Asp Val Leu Pro Trp Ile Ile

Cys Lys Gly Asp Trp Ser Arg Tyr Asn Glu Ala Arg Pro Pro Asn Asn 355 360 365

Gly Gln Lys Cys Thr Glu Ser Pro Ser Asp Glu Asp Tyr Ile Lys Gln 370 375 380

Phe Gln Glu Ala Arg Glu Tyr 385 390

<210> 403

<211> 538

<212> PRT

<213> Homo sapiens

<400> 403

Val His Ser His Gly Asp Lys Asp Ser Gln Thr Cys Ile Arg Val Ser

1 5 10 15

Ala Ser Pro Asp Pro Arg Pro Leu Lys Glu Glu Glu Glu Ala Pro Leu
20 25 30

Leu Pro Arg Thr His Leu Gln Ala Glu Pro His Gln His Gly Cys Trp
35 40 45

Thr Val Thr Glu Pro Ala Ala Met Thr Pro Gly Asn Ala Thr Pro Pro 50 55 60

Arg Thr Pro Glu Val Thr Pro Leu Arg Leu Glu Leu Gln Lys Leu Pro 65 70 75 80

Gly Leu Ala Asn Thr Thr Leu Ser Thr Pro Asn Pro Asp Thr Gln Ala 85 90 95

Ser Ala Ser Pro Asp Pro Arg Pro Leu Arg Glu Glu Glu Glu Ala Arg 100 105 110

Leu Leu Pro Arg Thr His Leu Gln Ala Glu Leu His Gln His Gly Cys
115 120 125

Trp Thr Val Thr Glu Pro Ala Ala Leu Thr Pro Gly Asn Ala Thr Pro
130 135 140

Pro Arg Thr Gln Glu Val Thr Pro Leu Leu Glu Leu Gln Lys Leu 145 150 155 160

Pro Glu Leu Val His Ala Thr Leu Ser Thr Pro Asn Pro Asp Asn Gln

| | | | | 165 | | | | | 170 | | | | | 175 | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Val | Thr | Ile | Lys 180 | Val | Val | Glu | Asp | Pro 185 | Gln | Ala | Glu | Val | Ser 190 | Ile | Asp |
| Leu | Leu | Ala 195 | Glu | Pro | Ser | Asn | Pro 200 | Pro | Pro | Gln | Asp | Thr 205 | Leu | Ser | Trp |
| Leu | Pro 210 | Ala | Leu | Trp | Ser | Phe 215 | Leu | Trp | Gly | Asp | Tyr 220 | Lys | Gly | Glu | Glu |
| Lys 225 | Asp | Arg | Ala | Pro | Glÿ 230 | Glu | Lys | Gly | Glu | Glu 235 | Lys | Glu | Glu | Asp | Glu 240 |
| Asp | Tyr | Pro | Ser | Glu 245 | Asp | Ile | Glu | Gly | Glu 250 | Asp | Gln | Glu | Asp | Lys 255 | Glu |
| Glu | Asp | Glu | Glu 260 | Glu | Gln | Ala | Leu | Trp 265 | Phe | Asn | Gly | Thr | Thr 270 | Asp | Asn |
| Trp | Asp | Gln 275 | Gly | Trp | Leu | Ala | Pro 280 | Gly | Asp | Trp | Val | Phe 285 | Lys | Asp | Ser |
| Val | Ser 290 | Tyr | Asp | Tyr | Glu | Pro 295 | Gln | Lys | Glu | Trp | Ser 300 | Pro | Trp | Ser | Pro |
| Cys 305 | Ser | Gly | Asn | Cys | Ser 310 | Thr | Gly | Lys | Gln | Gln 315 | Arg | Thr | Arg | Pro | Cys 320 |
| Gly | Tyr | Gly | Cys | Thr 325 | Ala | Thr | Glu | Thr | Arg 330 | Thr | Cys | Asp | Leu | Pro 335 | Ser |
| Cys | Pro | Gly | Thr 340 | Glu | Asp | Lys | Asp | Thr 345 | Leu | Gly | Leu | Pro | Ser 350 | Glu | Glu |
| Trp | Lys | Leu 355 | Leu | Ala | Arg | Asn | Ala 360 | Thr | Asp | Met | His | Asp 365 | Gln | Asp | Val |
| Asp | Ser 370 | Cys | Glu | Lys | Trp | Leu 375 | Asn | Cys | Lys | Ser | Asp 380 | Phe | Leu | Ile | Lys |
| Tyr 385 | Leu | Ser | Gln | Met | Leu 390 | Arg | Asp | Leu | Pro | Ser 395 | Cys | Pro | Cys | Ala | Tyr 400 |
| Pro | Leu | Glu | Ala | Met 405 | Asp | Ser | Pro | Val | Ser 410 | Leu | Gln | Asp | Glu | His 415 | Gln |

Gly Arg Ser Phe Arg Trp Arg Asp Ala Ser Gly Pro Arg Glu Arg Leu

420 425 430

Asp Ile Tyr Gln Pro Thr Ala Arg Phe Cys Leu Arg Ser Met Leu Ser 435 440 445

Gly Glu Ser Ser Thr Leu Ala Ala Gln His Cys Cys Tyr Asp Glu Asp 450 455 460

Ser Arg Leu Leu Thr Arg Gly Lys Gly Ala Gly Met Pro Asn Leu Ile 465 470 475 480

Ser Thr Asp Phe Ser Pro Lys Leu His Phe Lys Phe Asp Thr Thr Pro
485 490 495

Trp Ile Leu Cys Lys Gly Asp Trp Ser Arg Leu His Ala Val Leu Pro 500 505 510

Pro Asn Asn Gly Arg Ala Cys Thr Asp Asn Pro Leu Glu Glu Glu Tyr 515 520 525

Leu Ala Gln Leu Gln Glu Ala Lys Glu Tyr 530 535

<210> 404

<211> 151

<212> PRT

<213> Homo sapiens

<400> 404

Lys Val Met Asn Asp Leu Pro Ser Cys Pro Cys Ser Tyr Pro Thr Glu
1 5 10 15

Val Ala Tyr Ser Thr Ala Asp Ile Phe Asp Arg Ile Lys Arg Lys Asp
20 25 30

Phe Arg Trp Lys Asp Ala Ser Gly Pro Lys Glu Lys Leu Glu Ile Tyr 35 40 45

Lys Pro Thr Ala Arg Tyr Cys Ile Arg Ser Met Leu Ser Leu Glu Ser 50 55 60

Thr Thr Leu Ala Ala Gln His Cys Cys Tyr Gly Asp Asn Met Gln Leu 65 70 75 80

Ile Thr Arg Gly Lys Gly Ala Gly Thr Pro Asn Leu Ile Ser Thr Glu 85 90 95

Phe Ser Ala Glu Leu His Tyr Lys Val Asp Val Leu Pro Trp Ile Ile 100 105 110 Cys Lys Gly Asp Trp Ser Arg Tyr Asn Glu Ala Arg Pro Pro Asn Asn 115 Gly Gln Lys Cys Thr Glu Ser Pro Ser Asp Glu Asp Tyr Ile Lys Gln 130 135 140 Phe Gln Glu Ala Arg Glu Tyr 145 150 <210> 405 <211> 56 <212> PRT <213> Homo sapiens <400> 405 Val Gly Ser Asp Thr Thr Ser Glu Thr Ser Phe Ser Leu Ser Lys Glu 5 10 Ala Pro Arg Glu His Leu Asp His Gln Ala Ala His Gln Pro Phe Pro 20 25 Arg Pro Arg Phe Arg Gln Glu Thr Gly His Pro Ser Leu Gln Arg Asp 35 Phe Pro Arg Ser Phe Leu Leu Asp 50 55 <210> 406 <211> 42 <212> PRT <213> Homo sapiens <400> 406 Gly Asp Trp Ser Leu Trp Ser Val Cys Ser Val Thr Cys Gly Asn Gly 10

Asn Gln Lys Arg Thr Arg Ser Cys Gly Tyr Ala Cys Thr Ala Thr Glu

25

20

35

Ser Arg Thr Cys Asp Arg Pro Asn Cys Pro

```
<210> 407
<211> 50
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Thrombospondin
      type 1 domain sequence
<400> 407
Gly Glu Trp Ser Glu Trp Ser Pro Cys Ser Val Thr Cys Gly Gly Gly
                                      10
Val Gln Thr Arg Thr Arg Cys Cys Asn Pro Pro Pro Asn Gly Gly Gly
             20
                                  25
Pro Cys Thr Gly Pro Asp Thr Glu Thr Arg Ala Cys Asn Glu Gln Pro
                             40
Cys Pro
     50
<210> 408
<211> 41
<212> PRT
<213> Homo sapiens
<400> 408
Gly Asp Trp Ser Leu Trp Ser Val Cys Ser Val Thr Cys Gly Asn Gly
 1
                  5
                                      10
Asn Gln Lys Arg Thr Arg Ser Cys Gly Tyr Ala Cys Thr Ala Thr Glu
             20
                                  25
Ser Arg Thr Cys Asp Arg Pro Asn Cys
         35
                              40
<210> 409
<211> 48
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: Thrombospondin
```

type 1 domain sequence

<400> 409

Ser Pro Trp Ser Glu Trp Ser Pro Cys Ser Val Thr Cys Gly Lys Gly
1 5 10 15

Ile Arg Thr Arg Gln Arg Thr Cys Asn Ser Pro Ala Gly Gly Lys Pro
20 25 30

Cys Thr Gly Asp Ala Gln Glu Thr Glu Ala Cys Met Met Asp Pro Cys
35 40 45

<210> 410

<211> 460

<212> PRT

<213> Homo sapiens

<400> 410

Met Ala Gly Tyr Leu Ser Glu Ser Asp Phe Val Met Val Glu Glu Gly 1 5 10 15

Phe Ser Thr Arg Asp Leu Leu Lys Glu Leu Thr Leu Gly Ala Ser Gln 20 25 30

Ala Thr Thr Asp Glu Val Ala Ala Phe Phe Val Ala Asp Leu Gly Ala 35 40 45

Ile Val Arg Lys His Phe Cys Phe Leu Lys Cys Leu Pro Arg Val Arg 50 55 60

Pro Phe Tyr Ala Val Lys Cys Asn Ser Ser Pro Gly Val Leu Lys Val 65 70 75 80

Leu Ala Gln Leu Gly Leu Gly Phe Ser Cys Ala Asn Lys Ala Glu Met 85 90 95

Glu Leu Val Gln His Ile Gly Ile Pro Ala Ser Lys Ile Ile Cys Ala 100 105 110

Asn Pro Cys Lys Gln Ile Ala Gln Ile Lys Tyr Ala Ala Lys His Gly
115 120 125

Ile Gln Leu Leu Ser Phe Asp Asn Glu Met Glu Leu Ala Lys Val Val 130 135 140

Lys Ser His Pro Ser Ala Lys Met Val Leu Cys Ile Ala Thr Asp Asp

| 145 150 | 155 | 160 |
|---------|-----|-----|
|---------|-----|-----|

| Ser His Ser Leu | Ser Cys Leu | Ser Leu Lys Phe | Gly Val Ser Leu Lys |
|-----------------|-------------|-----------------|---------------------|
| | 165 | 170 | 175 |

- Ser Cys Arg His Leu Leu Glu Asn Ala Lys Lys His His Val Glu Val 180 185 190
- Val Gly Val Ser Phe His Ile Gly Ser Gly Cys Pro Asp Pro Gln Ala 195 200 205
- Tyr Ala Gln Ser Ile Ala Asp Ala Arg Leu Val Phe Glu Met Gly Thr 210 215 220
- Glu Leu Gly His Lys Met His Val Leu Asp Leu Gly Gly Gly Phe Pro 225 230 235 240
- Gly Thr Glu Gly Ala Lys Val Arg Phe Glu Glu Ile Ala Ser Val Ile 245 250 255
- Asn Ser Ala Leu Asp Leu Tyr Phe Pro Glu Gly Cys Gly Val Asp Ile 260 265 270
- Phe Ala Glu Leu Gly Arg Tyr Tyr Val Thr Ser Ala Phe Thr Val Ala 275 280 285
- Val Ser Ile Ile Ala Lys Lys Glu Val Leu Leu Asp Gln Pro Gly Arg 290 295 300
- Glu Glu Glu Asn Gly Ser Thr Ser Lys Thr Ile Val Tyr His Leu Asp 305 310 315 320
- Glu Gly Val Tyr Gly Ile Phe Asn Ser Val Leu Phe Asp Asn Ile Cys 325 330 335
- Pro Thr Pro Ile Leu Gln Lys Lys Pro Ser Thr Glu Gln Pro Leu Tyr 340 345 350
- Ser Ser Ser Leu Trp Gly Pro Ala Val Asp Gly Cys Asp Cys Val Ala 355 360 365
- Glu Gly Leu Trp Leu Pro Gln Leu His Val Gly Asp Trp Leu Val Phe 370 380
- Asp Asn Met Gly Ala Tyr Thr Val Gly Met Gly Ser Pro Phe Trp Gly 385 390 395 400
- Thr Gln Ala Cys His Ile Thr Tyr Ala Met Ser Arg Val Ala Trp Glu

405 410 415

Ala Leu Arg Arg Gln Leu Met Ala Ala Glu Gln Glu Asp Asp Val Glu
420 425 430

Gly Val Cys Lys Pro Leu Ser Cys Gly Trp Glu Ile Thr Asp Thr Leu
435 440 445

Cys Val Gly Pro Val Phe Thr Pro Ala Ser Ile Met 450 455 460

<210> 411

<211> 480

<212> PRT

<213> Homo sapiens

<400> 411

Met Ala Gly Tyr Leu Ser Glu Ser Asp Phe Val Met Val Glu Glu Gly 1 5 10 15

Phe Ser Thr Arg Asp Leu Leu Lys Glu Leu Thr Leu Gly Ala Ser Gln 20 25 30

Ala Thr Thr Asp Glu Val Ala Ala Phe Phe Val Ala Asp Leu Gly Ala 35 40 45

Ile Val Arg Lys His Phe Cys Phe Leu Lys Cys Leu Pro Arg Val Arg 50 55 60

Pro Phe Tyr Ala Val Lys Cys Asn Ser Ser Pro Gly Val Leu Lys Val 65 70 75 80

Leu Ala Gln Leu Gly Leu Gly Phe Ser Cys Ala Asn Lys Ala Glu Met 85 90 95

Glu Leu Val Gln His Ile Gly Ile Pro Ala Ser Lys Ile Ile Cys Ala 100 105 110

Asn Pro Cys Lys Gln Ile Ala Gln Ile Lys Tyr Ala Ala Lys His Gly
115 120 125

Ile Gln Leu Leu Ser Phe Asp Asn Glu Met Glu Leu Ala Lys Val Val 130 135 140

Lys Ser His Pro Ser Ala Lys Met Val Leu Cys Ile Ala Thr Asp Asp 145 150 155 160

| Ser | His | Ser | Leu | Ser 165 | Cys | Leu | Ser | Leu | Lys 170 | Phe | Gly | Val | Ser | Leu 175 | Lys |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser | Cys | Arg | His 180 | Leu | Leu | Glu | Àsn | Ala 185 | Lys | Lys | His | His | Val 190 | Glu | Val |
| Val | Gly | Val 195 | Ser | Phe | His | Ile | Gly 200 | Ser | Gly | Cys | Pro | Asp 205 | Pro | Gln | Ala |
| Tyr | Ala 210 | Gln | Ser | Ile | Ala | Asp 215 | Ala | Arg | Leu | Val | Phe 220 | Glu | Met | Gly | Thr |
| Glu 225 | Leu | Gly | His | Lys | Met 230 | His | Val | Leu | Asp | Leu 235 | Gly | Gly | Gly | Phe | Pro 240 |
| Gly | Thr | Glu | Gly | Ala 245 | Lys | Val | Arg | Phe | Glu 250 | Glu | Ile | Ala | Ser | Val 255 | Ile |
| Asn | Ser | Ala | Leu 260 | Asp | Leu | Tyr | Phe | Pro 265 | Glu | Gly | Cys | Gly | Val 270 | Asp | Ile |
| Phe | Ala | Glu 275 | Leu | Gly | Arg | Tyr | Tyr 280 | Val | Thr | Ser | Ala | Phe 285 | Thr | Val | Ala |
| Val | Ser 290 | Ile | Ile | Ala | Lys | Lys 295 | Glu | Val | Leu | Leu | Asp 300 | Gln | Pro | Gly | Arg |
| Glu 305 | Ala | Pro | Leu | Pro | Pro 310 | Pro | His | Ile | Ala | Thr 315 | Cys | Ala | Ala | Ser | Glu 320 |
| Pro | Ser | Pro | Pro | Ala 325 | Glu | Glu | Asn | Gly | Ser 330 | Thr | Ser | Lys | Thr | Ile 335 | Val |
| Tyr | His | Leu | Asp 340 | Glu | Gly | Val | Tyr | Gly 345 | Ile | Phe | Asn | Ser | Val 350 | Leu | Phe |
| Asp | Asn | Ile 355 | Cys | Pro | Thr | Pro | Ile 360 | Leu | Gln | Lys | Lys | Pro 365 | Ser | Thr | Glu |
| Gln | Pro 370 | Leu | Tyr | Ser | Ser | Ser 375 | Leu | Trp | Gly | Pro | Ala 380 | Val | Asp | Gly | Cys |
| Asp 385 | Cys | Val | Ala | Glu | Gly 390 | Leu | Trp | Leu | Pro | Gln 395 | Leu | His | Val | Gly | Asp 400 |
| Trp | Leu | Val | Phe | Asp | Asn | Met | Gly | Ala | Tyr 410 | Thr | Val | Gly | Met | Gly 415 | Ser |

Pro Phe Trp Gly Thr Gln Ala Cys His Ile Thr Tyr Ala Met Ser Arg 420 425 430

Val Ala Trp Glu Ala Leu Arg Arg Gln Leu Met Ala Ala Glu Gln Glu
435 440 445

Asp Asp Val Glu Gly Val Cys Lys Pro Leu Ser Cys Gly Trp Glu Ile 450 455 460

Thr Asp Thr Leu Cys Val Gly Pro Val Phe Thr Pro Ala Ser Ile Met 465 470 475 480

<210> 412

<211> 365

<212> PRT

<213> Homo sapiens

<400> 412

Met Glu Leu Val Gln His Ile Gly Ile Pro Ala Ser Lys Ile Ile Cys
1 5 10 15

Ala Asn Pro Cys Lys Gln Ile Ala Gln Ile Lys Tyr Ala Ala Lys His
20 25 30

Gly Ile Gln Leu Leu Ser Phe Asp Asn Glu Met Glu Leu Ala Lys Val 35 40 45

Val Lys Ser His Pro Ser Ala Lys Met Val Leu Cys Ile Ala Thr Asp 50 55 60

Asp Ser His Ser Leu Ser Cys Leu Ser Leu Lys Phe Gly Val Ser Leu 65 70 75 80

Lys Ser Cys Arg His Leu Leu Glu Asn Ala Lys Lys His His Val Glu 85 90 95

Val Val Gly Val Ser Phe His Ile Gly Ser Gly Cys Pro Asp Pro Gln 100 105 110

Ala Tyr Ala Gln Ser Ile Ala Asp Ala Arg Leu Val Phe Glu Met Gly
115 120 125

Thr Glu Leu Gly His Lys Met His Val Leu Asp Leu Gly Gly Phe 130 135 140

| Pro Gly 145 | Thr (| Glu (| Gly | Ala 150 | Lys | Val | Arg | Phe | Glu 155 | Glu | Ile | Ala | Ser | Val 160 |
|----------------|--------------|--------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile Asn | Ser A | | Leu 165 | Asp | Leu | Tyr | Phe | Pro 170 | Glu | Gly | Cys | Gly | Val 175 | Asp |
| Ile Phe | | Glu : 180 | Leu | Gly | Arg | Tyr | Tyr 185 | Val | Thr | Ser | Ala | Phe 190 | Thr | Val |
| Ala Val | Ser 1 | Ile : | Ile | Ala | Lys | Lys 200 | Glu | Val | Leu | Leu | Asp 205 | Gln | Pro | Gly |
| Arg Glu 210 | Glu (| Glu <i>i</i> | Asn | Gly | Ser 215 | Thr | Ser | Lys | Thr | Ile 220 | Val | Tyr | His | Leu |
| Asp Glu 225 | _ | | | 230 | | | | | 235 | | | | | 240 |
| Cys Pro | | : | 245 | | | | | 250 | | | | | 255 | |
| Tyr Ser | | Ser : 260 | Leu | Trp | Gly | Pro | Ala 265 | Val | Asp | Gly | Cys | Asp 270 | Cys | Val |
| Ala Glu | Gly I 275 | Leu ' | Trp | Leu | Pro | Gln 280 | Leu | His | Val | Gly | Asp 285 | Trp | Leu | Val |
| Phe Asp 290 | Asn N | Met (| Gly | Ala | Tyr 295 | Thr | Val | Gly | Met | Gly 300 | Ser | Pro | Phe | Trp |
| Gly Thr 305 | Gln A | Ala (| Cys | His 310 | Ile | Thr | Tyr | Ala | Met 315 | Ser | Arg | Val | Ala | Trp 320 |
| Glu Ala | Leu A | | Arg 325 | Gln | Leu | Met | Ala | Ala 330 | Glu | Gln | Glu | Asp | Asp 335 | Val |
| Glu Gly | | Cys : | Lys | Pro | Leu | Ser | Cys 345 | Gly | Trp | Glu | Ile | Thr 350 | Asp | Thr |
| Leu Cys | Val (| Gly 1 | Pro | Val | Phe | Thr 360 | Pro | Ala | Ser | Ile | Met 365 | | | |

<210> 413

<211> 362

<212> PRT

<213> Homo sapiens

| <400 |)> 41 | L3 | | | | | | | | | | | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Met 1 | Ala | Gly | Tyr | Leu 5 | Ser | Glu | Ser | Asp | Phe 10 | Val | Met | Val | Glu | Glu 15 | Gly |
| Phe | Ser | Thr | Arg 20 | Asp | Leu | Leu | Lys | Glu 25 | Leu | Thr | Leu | Gly | Ala 30 | Ser | Gln |
| Ala | Thr | Thr 35 | Asp | Glu | Val | Ala | Ala 40 | Phe | Phe | Val | Ala | Asp 45 | Leu | Gly | Ala |
| Ile | Val 50 | Arg | Lys | His | Phe | Cys 55 | Phe | Leu | Lys | Cys | Leu 60 | Pro | Arg | Val | Arg |
| Pro 65 | Phe | Tyr | Ala | Val | Lys 70 | Cys | Asn | Ser | Ser | Pro 75 | Gly | Val | Leu | Lys | Val 80 |
| Leu | Ala | Gln | Leu | Gly 85 | Leu | Gly | Phe | Ser | Cys 90 | Ala | Asn | Lys | Ala | Glu 95 | Met |
| Glu | Leu | Val | Gln 100 | His | Ile | Gly | Ile | Pro 105 | Ala | Ser | Lys | Ile | Ile 110 | Cys | Ala |
| Asn | Pro | Cys 115 | Lys | Gln | Ile | Ala | Gln 120 | Ile | Lys | Tyr | Ala | Ala 125 | Lys | His | Gly |
| Ile | Gln 130 | Leu | Leu | Ser | Phe | Asp 135 | Asn | Glu | Met | Glu | Leu 140 | Ala | Lys | Val | Val |
| Lys 145 | Ser | His | Pro | Ser | Ala 150 | Lys | Met | Val | Leu | Cys 155 | Ile | Ala | Thr | Asp | Asp 160 |
| Ser | His | Ser | Leu | Ser 165 | Cys | Leu | Ser | Leu | Lys 170 | Phe | Gly | Val | Ser | Leu 175 | Lys |
| Ser | Cys | Arg | His 180 | Leu | Leu | Glu | Asn | Ala 185 | Lys | Lys | His | His | Val 190 | Glu | Val |
| Val | Gly | Val 195 | Ser | Phe | His | Ile | Gly 200 | Ser | Gly | Cys | Pro | Asp 205 | Pro | Gln | Ala |
| Tyr | Ala 210 | Gln | Ser | Ile | Ala | Asp 215 | Ala | Arg | Leu | Val | Phe 220 | Glu | Met | Gly | Thr |

Glu Leu Gly His Lys Met His Val Leu Asp Leu Gly Gly Phe Pro

Gly Thr Glu Gly Ala Lys Val Arg Phe Glu Glu Ile Ala Ser Val Ile

235

230

Asn Ser Ala Leu Asp Leu Tyr Phe Pro Glu Gly Cys Gly Val Asp Ile 260 265 270

Phe Ala Glu Leu Gly Arg Tyr Tyr Val Thr Ser Ala Phe Thr Val Ala 275 280 285

Val Ser Ile Ile Ala Lys Lys Glu Val Leu Leu Asp Gln Pro Gly Arg 290 295 300

Glu Glu Glu Asn Gly Ser Thr Ser Lys Thr Ile Val Tyr His Leu Asp 305 310 315 320

Glu Gly Val Tyr Gly Ile Phe Asn Ser Val Leu Phe Asp Asn Ile Cys 325 330 335

Pro Thr Pro Ile Leu Gln Lys Ser Lys Asn His Ser Pro Cys Tyr Met 340 345 350

Ser Leu Glu Ser Ile His Phe Ile Ala Val 355 360

<210> 414

<211> 374

<212> PRT

<213> Homo sapiens

<400> 414

Met Ala Gly Tyr Leu Ser Glu Ser Asp Phe Val Met Val Glu Glu Gly 1 5 10 15

Phe Ser Thr Arg Asp Leu Leu Lys Glu Leu Thr Leu Gly Ala Ser Gln 20 25 30

Ala Thr Thr Asp Glu Val Ala Ala Phe Phe Val Ala Asp Leu Gly Ala
35 40 45

Ile Val Arg Lys His Phe Cys Phe Leu Lys Cys Leu Pro Arg Val Arg 50 55 60

Pro Phe Tyr Ala Val Lys Cys Asn Ser Ser Pro Gly Val Leu Lys Val 65 70 75 80

Leu Ala Gln Leu Gly Leu Gly Phe Ser Cys Ala Asn Lys Ala Glu Met 85 90 95

| Glu | Leu | Val | Gln 100 | His | Ile | Gly | Ile | Pro 105 | Ala | Ser | Lys | Ile | Ile 110 | Cys | Ala |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Asn | Pro | Cys 115 | Lys | Gln | Ile | Ala | Gln 120 | Ile | Lys | Tyr | Ala | Ala 125 | Lys | His | Gly |
| Ile | Gln 130 | Leu | Leu | Ser | Phe | Asp 135 | Asn | Glu | Met | Glu | Leu 140 | Ala | Lys | Val | Val |
| Lys 145 | Ser | His | Pro | Ser | Ala 150 | Lys | Phe | Val | Gln | Gln 155 | Arg | Gly | Thr | Ala | Cys 160 |
| Leu | Ile | Arg | Met | Val 165 | Leu | Cys | Ile | Ala | Thr 170 | Asp | Asp | Ser | His | Ser 175 | Leu |
| Ser | Cys | Leu | Ser 180 | Leu | Lys | Phe | Gly | Val 185 | Ser | Leu | Lys | Ser | Cys 190 | Arg | His |
| Leu | Leu | Glu 195 | Asn | Ala | Lys | Lys | His 200 | His | Val | Glu | Val | Val 205 | Gly | Val | Ser |
| Phe | His 210 | Ile | Gly | Ser | Gly | Cys 215 | Pro | Asp | Pro | Gln | Ala 220 | Tyr | Ala | Gln | Ser |
| Ile 225 | Ala | Asp | Ala | Arg | Leu 230 | Val | Phe | Glu | Met | Gly 235 | Thr | Glu | Leu | Gly | His 240 |
| Lys | Met | His | Val | Leu 245 | Asp · | Leu | Gly | Gly | Gly 250 | Phe | Pro | Gly | Thr | Glu 255 | Gly |
| Ala | Lys | Val | Arg 260 | Phe | Glu | Glu | Ile | Ala 265 | Ser | Val | Ile | Asn | Ser 270 | Ala | Leu |
| Asp | Leu | Tyr 275 | Phe | Pro | Glu | Gly | Cys 280 | Gly | Val | Asp | Ile | Phe 285 | Ala | Glu | Leu |
| Gly | Arg 290 | Tyr | Tyr | Val | Thr | Ser 295 | Ala | Phe | Thr | Val | Ala 300 | Val | Ser | Ile | Ile |
| Ala 305 | Lys | Lys | Glu | Val | Leu 310 | Leu | Asp | Gln | Pro | Gly 315 | Arg | Glu | Glu | Glu | Asn 320 |
| Gly | Ser | Thr | Ser | Lys 325 | Thr | Ile | Val | Tyr | His 330 | Leu | Asp | Glu | Gly | Val 335 | Tyr |
| Gly | Ile | Phe | Asn | Ser | Val | Leu | Phe | Asp | Asn | Ile | Cys | Pro | Thr 350 | Pro | Ile |

Leu Gln Lys Ser Lys Asn His Ser Pro Cys Tyr Met Ser Leu Glu Ser 355 360 365

Ile His Phe Ile Ala Val 370

<210> 415

<211> 237

<212> PRT

<213> Homo sapiens

<400> 415

Asp Leu Gly Ala Ile Val Arg Lys His Phe Cys Phe Leu Lys Cys Leu
1 5 10 15

Pro Arg Val Arg Pro Phe Tyr Ala Val Lys Cys Asn Ser Ser Pro Gly
20 25 30

Val Leu Lys Val Leu Ala Gln Leu Gly Leu Gly Phe Ser Cys Ala Asn 35 40 45

Lys Ala Glu Met Glu Leu Val Gln His Ile Gly Ile Pro Ala Ser Lys
50 55 60

Ile Ile Cys Ala Asn Pro Cys Lys Gln Ile Ala Gln Ile Lys Tyr Ala 65 70 75 80

Ala Lys His Gly Ile Gln Leu Leu Ser Phe Asp Asn Glu Met Glu Leu 85 90 95

Ala Lys Val Val Lys Ser His Pro Ser Ala Lys Met Val Leu Cys Ile 100 105 110

157

Ala Thr Asp Asp Ser His Ser Leu Ser Cys Leu Ser Leu Lys Phe Gly
115 120 125

Val Ser Leu Lys Ser Cys Arg His Leu Leu Glu Asn Ala Lys Lys His 130 135 140

His Val Glu Val Val Gly Val Ser Phe His Ile Gly Ser Gly Cys Pro 145 150 155 160

Asp Pro Gln Ala Tyr Ala Gln Ser Ile Ala Asp Ala Arg Leu Val Phe 165 170 175

Glu Met Gly Thr Glu Leu Gly His Lys Met His Val Leu Asp Leu Gly
180 185 190

Gly Gly Phe Pro Gly Thr Glu Gly Ala Lys Val Arg Phe Glu Glu Ile 195 200 205 Ala Ser Val Ile Asn Ser Ala Leu Asp Leu Tyr Phe Pro Glu Gly Cys 210 215 220 Gly Val Asp Ile Phe Ala Glu Leu Gly Arg Tyr Tyr Val 225 230 235 <210> 416 <211> 244 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Pyridoxal-dependent decarboxylase domain sequence <400> 416 Asp Leu Gly Leu Ile Val Arg Arg Ile His Ala Leu Trp Gln Ala Phe 10 Leu Pro Arg Ile Gln Pro Phe Tyr Ala Val Lys Ala Asn Ser Asp Pro 20 25 Ala Val Leu Arg Leu Leu Ala Glu Leu Gly Thr Gly Phe Asp Cys Ala 40 Ser Lys Gly Glu Leu Glu Arg Val Leu Ala Ala Gly Val Pro Pro Glu Arg Ile Ile Phe Ala Asn Pro Cys Lys Asp Arg Ser Glu Leu Arg Tyr 70 75 Ala Leu Glu His Gly Val Val Cys Val Thr Val Asp Asn Val Glu Glu 85 90 95 Leu Glu Lys Leu Ala Arg Leu Ala Pro Glu Ala Arg Leu Leu Leu Arg 100 105 110 Val Lys Pro Asp Val Asp Ala His Ala His Cys Tyr Leu Ser Thr Gly 115 120 125

135

130

Gln Asp Ser Lys Phe Gly Ala Asp Leu Glu Glu Ala Glu Ala Leu Leu

140

Lys Ala Ala Lys Glu Leu Gly Leu Asn Val Val Gly Val His Phe His Val Gly Ser Gly Cys Thr Asp Ala Glu Ala Phe Val Lys Ala Ala Arg Asp Ala Arg Asn Val Phe Asp Gln Gly Ala Asp Glu Leu Gly Phe Glu Leu Lys Ile Leu Asp Leu Gly Gly Gly Phe Gly Val Asp Tyr Thr Gly Ala Glu Asp Phe Glu Glu Tyr Ala Glu Val Ile Asn Ala Ala Leu Glu Glu Val Phe Pro His Asp Pro His Pro Thr Ile Ile Ala Glu Pro Gly Arg Tyr Ile Val <210> 417 <211> 112 <212> PRT <213> Homo sapiens <400> 417 Val Ala Val Ser Ile Ile Ala Lys Lys Glu Val Leu Leu Asp Gln Pro Gly Arg Glu Glu Asn Gly Ser Thr Ser Lys Thr Ile Val Tyr His Leu Asp Glu Gly Val Tyr Gly Ile Phe Asn Ser Val Leu Phe Asp Asn Ile Cys Pro Thr Pro Ile Leu Gln Lys Lys Pro Ser Thr Glu Gln Pro Leu Tyr Ser Ser Leu Trp Gly Pro Ala Val Asp Gly Cys Asp Cys Val Ala Glu Gly Leu Trp Leu Pro Gln Leu His Val Gly Asp Trp Leu Val Phe Asp Asn Met Gly Ala Tyr Thr Val Gly Met Gly Ser Pro Phe

<210> 418 <211> 107 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Pyridoxal-dependent decarboxylase domain sequence <400> 418 Thr Leu Val Ser Asn Val Ile Ala Lys Lys Thr Val Pro Ser Asp Asp 10 Glu Asp Gly Lys Asp Asp Thr Arg Met Tyr Tyr Val Asn Asp Gly Gly 20 25 Tyr Ser Ser Phe Ile Arg Pro Leu Leu Tyr His Ala His Pro His Ala 35 40 Leu Leu Arg Arg Ser Leu Asp Glu Glu Pro Pro Arg Lys Ser Ser 55 Ile Trp Gly Pro Thr Cys Asp Ser Leu Asp Lys Ile Ile Lys Asp Arg 70 75 Leu Leu Pro Glu Leu Asp Val Gly Asp Trp Leu Ala Phe Phe Asp Thr 90 Gly Ala Tyr Thr Glu Ala Met Ala Ser Asn Phe 100 105 <210> 419

15

<211> 467

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Pyridoxal-dependent decarboxylase domain sequence

<400> 419

Phe Tyr Val Tyr Asp Leu Gly Leu His Ile Val Arg Arg Ile His Ala

- Leu Trp Lys Ala Phe Leu Pro Arg Gly Gln Tyr Asn Ser Val Val Lys
 20 25 30
- Pro Phe Tyr Ala Val Lys Ala Asn Ser Asp Pro Ala Val Leu Arg Leu
 35 40 45
- Leu Ala Glu Leu Gly Thr His Ser Leu Gly Phe Asp Cys Ala Ser Lys 50 55 60
- Gly Glu Leu Glu Arg Val Leu Ala Ala Tyr Leu Ala Gly Val Ser Pro 65 70 75 80
- Glu Arg Ile Ile Phe Ala Asn Pro Cys Lys Ser Arg Ser Glu Leu Arg 85 90 95
- Tyr Ala Leu Glu His Arg Lys Met Gly Gly Val Val Cys Val Thr Val
 100 105 110
- Asp Asn Val Glu Glu Leu Glu Lys Ile Ala Lys Leu Ala Pro Glu Ala 115 120 125
- Gly Val Lys Pro Arg Leu Leu Leu Arg Val Lys Pro Asp Val Asp Ala 130 135 140
- His Ala His Cys Arg Leu Ser Thr Gly Gln Glu Asp Ser Lys Phe Gly
 145 150 155 160
- Ala Asp Leu Glu Asp Gly Glu Asp Ala Glu Ala Leu Leu Lys Ala Ala 165 170 175
- Lys Glu Leu Gly Asn Leu Asn Val Val Gly Val His Phe His Val Gly
 180 185 190
- Ser Gly Ile Ser Asp Leu Glu Ala Phe Val Lys Ala Val Arg Asp Ala 195 200 205
- Arg Asn Val Phe Asp Gln Gly Ala Asp Glu Leu Gly Phe Lys Thr Ile 210 215 220
- Asp Leu Lys Ile Leu Asp Ile Gly Gly Gly Phe Gly Val Asp Tyr Thr 225 230 235 240
- Gly Thr Arg Ser Gln Ser Asp Met Ser Val Ala Glu Asp Phe Glu Glu 245 250 255
- Ile Ala Glu Val Ile Asn Ala Ala Leu Glu Glu Leu Phe Pro His Ala

| GLY | Tyr | GIY | Asp | Pro | GLY | Pro | Thr | TTE | TT6 | Ala | Glu | Pro | GTA | Arg | Tyr |
|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|
| | | 275 | | | | | 280 | | | | | 285 | | | |

Ile Val Ala Ala Ala Gly Thr Leu Val Ser Asn Val Ile Ala Lys Lys 290 295 300

Glu Val Pro Ser Asp Asp Ala Asp Thr Thr Ser Asp Ser Leu Arg Glu 305 310 315 320

Glu Ser Lys Asp Asp Thr Arg Met Tyr Tyr Val Asn Asp Gly Gly Tyr 325 330 335

Gly Ser Phe Ile Arg Pro Leu Leu Tyr His Ala His Pro Glu Ala Leu 340 345 350

Leu Leu Arg Arg Gly Gly Glu Val Gln Tyr Gln Asp Ala Glu Thr Glu 355 360 365

Arg Ala Ala Asp Lys Ser Leu Ser Asn Phe Ser Leu Phe Gln Ser Tyr 370 380

Pro Asp Ala Trp Gly Ile Asp Gln Leu Phe Pro Val Leu Pro Leu Arg 385 390 395 400

Ser Leu Asp Glu Glu Pro Lys Arg Lys Ser Ser Ile Val Gly Pro Thr \$405\$

Cys Asp Ser Asp Gly Lys Leu Asp Lys Ile Ile Lys Asp Asp Gly Ile 420 425 430

Ala Glu Asp Arg Leu Leu Pro Glu Leu Lys Pro Val Gly Asp Trp Leu 435 440 445

Ala Phe Pro Asp Thr Gly Ala Tyr Thr Tyr Ala Met Ala Ser Asn Tyr 450 455 460

Asn Gly Phe 465

<210> 420

<211> 361

<212> PRT

<213> Homo sapiens

<400> 420

| Phe 1 | Phe | Val | Ala | Asp 5 | Leu | Gly | Ala | Ile | Val 10 | Arg | Lys | His | Phe | Cys 15 | Phe |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Lys | Cys | Leu 20 | Pro | Arg | Val | Arg | Pro 25 | Phe | Tyr | Ala | Val | Lys 30 | Cys | Asn |
| Ser | Ser | Pro 35 | Gly | Val | Leu | Lys | Val 40 | Leu | Ala | Gln | Leu | Gly 45 | Leu | Gly | Phe |
| Ser | Cys 50 | Ala | Asn | Lys | Ala | Glu 55 | Met | Glu | Leu | Val | Gln 60 | His | Ile | Gly | Ile |
| Pro 65 | Ala | Ser | Lys | Ile | Ile 70 | Cys | Ala | Asn | Pro | Cys 75 | Lys | Gln | Ile | Ala | Gln 80 |
| Ile | Lys | Tyr | Ala | Ala 85 | Lys | His | Gly | Ile | Gln 90 | Leu | Leu | Ser | Phe | Asp 95 | Asn |
| Glu | Met | Glu | Leu 100 | Ala | Lys | Val | Val | Lys 105 | Ser | His | Pro | Ser | Ala 110 | Lys | Met |
| Val | Leu | Cys 115 | Ile | Ala | Thr | Asp | Asp 120 | Ser | His | Ser | Leu | Ser 125 | Cys | Leu | Ser |
| Leu | Lys 130 | Phe | Gly | Val | Ser | Leu 135 | Lys | Ser | Cys | Arg | His 140 | Leu | Leu | Glu | Asn |
| Ala 145 | Lys | Lys | His | His | Val 150 | Glu | Val | Val | Gly | Val 155 | Ser | Phe | His | Ile | Gly 160 |
| Ser | Gly | Cys | Pro | Asp 165 | Pro | Gln | Ala | Tyr | Ala 170 | Gln | Ser | Ile | Ala | Asp 175 | Ala |
| Arg | Leu | Val | Phe 180 | Glu | Met | Gly | Thr | Glu 185 | Leu | Gly | His | Lys | Met 190 | His | Val |
| Leu | Asp | Leu 195 | Gly | Gly | Gly | Phe | Pro 200 | Gly | Thr | Glu | Gly | Ala 205 | Lys | Val | Arg |
| Phe | Glu 210 | Glu | Ile | Ala | Ser | Val 215 | Ile | Asn | Ser | Ala | Leu 220 | Asp | Leu | Tyr | Phe |
| Pro 225 | Glu | Gly | Cys | Gly | Val 230 | Asp | Ile | Phe | Ala | Glu 235 | Leu | Gly | Arg | Tyr | Tyr 240 |
| Val | Thr | Ser | Ala | Phe 245 | Thr | Val | Ala | Val | Ser 250 | Ile | Ile | Ala | Lys | Lys 255 | Glu |

Val Leu Leu Asp Gln Pro Gly Arg Glu Glu Asn Gly Ser Thr Ser Lys Thr Ile Val Tyr His Leu Asp Glu Gly Val Tyr Gly Ile Phe Asn Ser Val Leu Phe Asp Asn Ile Cys Pro Thr Pro Ile Leu Gln Lys Lys Pro Ser Thr Glu Gln Pro Leu Tyr Ser Ser Ser Leu Trp Gly Pro Ala Val Asp Gly Cys Asp Cys Val Ala Glu Gly Leu Trp Leu Pro Gln Leu His Val Gly Asp Trp Leu Val Phe Asp Asn Met Gly Ala Tyr Thr Val Gly Met Gly Ser Pro Phe Trp Gly Thr <210> 421 <211> 479 <212> PRT <213> Mus musculus <400> 421 Met Leu Gln Ile Thr Glu Trp Arg Phe Leu Ala Arg Asp Glu Gly Glu Ser Ala Val Ala Glu Asp Pro Thr Trp Gly Glu Asp Glu Glu Pro Leu Ala Cys Thr Thr Asp Ser Trp Ala Gln Gly Ser Val Pro Val Leu His Thr Pro Ala Pro Val Cys Val Glu Glu Gln Phe His Asn Glu Glu Pro Gly Asn Pro Asp Gln Phe Leu Leu Gly Ser Ser Trp Asp Lys Glu Ser Gln Lys Pro Thr Gln Pro Ser Glu Pro Ser Ala Glu Pro Lys Val Thr Pro Arg Pro Thr Ala Thr Leu Glu Ala Phe Glu Glu Ala Glu Pro Gly

| Leu Tyr Leu Cys Met Pro Gln Pro Asp Ala Ala Gly Asp Arg Leu 190 Arg Leu 185 Ala Ala Ala Ala Ala Arg Arg Arg Ala Ala Ala Ala Ala Pro Ala Pro Ala Pro Ala Pro Ala Pro Ala Ala Pro Ala Pro Ala Pro Ala Pro Ala Pro Ala Pro Ala Ala Pro Ala Ala Pro Ala Ala Pro Ala A | | | | | | | | | | | | | | | | |
|--|-----|-------|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|-----|------------|------------|
| 145 | | | Ser | Lys | Glu | Ser | | Arg | Ser | Thr | Ala | | Gly | Glu | Arg | Val |
| Leu Tyr Leu Cys Met 180 Pro Gln Pro Asp 185 Ala Ala Gly Asp Arg L 190 Leu Gln Ser Lys Gly Gln Leu His 200 Ser Ser Pro Ile Gly 205 Ser Gly 205 His Leu Gly Ala Leu Thr Pro Ala Glu Pro 215 Ser Ala Phe Gln Gly 225 Glu Val Leu Gly Glu Arg Pro Lys His Lys Thr Thr Thr Leu A 225 Ser Arg Leu Pro Arg His Trp 250 Thr Thr Thr Leu A 255 Asp Ser Ser Arg Leu Pro Arg His Trp 265 Ser Glu Ile Try Arg G 270 Pro Arg Arg Ser Gln Ala Gly Thr Ala Thr Ser Ala Cys Glu S 285 Ala Leu Ser Ser Arg Ala Pro 295 Ser Leu Asn Leu Gln Arg Cys Ala Thr Phe Arg Ala Leu Gly Pro A 315 Ser Leu Asn Leu Ala Gln Thr Ser Pro 330 Ser Phe Gly Ser Asn V 325 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Arg Asn Pro Ile Pa360 | _ | Ser P | Pro | Gln | Ser | | Leu | Lys | Gln | Pro | | Val | Val | Arg | Leu | Gln 160 |
| Leu Gln Ser Lys Gly Gln Leu His Ser Ser Pro Ile Gly Ser Green 195 | Ala | Ser G | Slu | Lys | | Ser | Ser | Phe | Gly | | His | Leu | Ser | Leu | Glu 175 | Asp |
| His Leu Gly Ala Leu Thr Pro Ala Glu Pro Ser Ala Phe Gln G 225 Asp Ser Ser Arg Leu Pro Arg His Trp Val Arg Pro Val Ala Glu Ile Tyr Arg G 260 Leu Ile Pro Asp Leu Glu Val His Pro Leu Glu Ile Tyr Arg G 270 Pro Arg Arg Ser Gln Ala Gly Thr Ala Thr Ser Ala Cys Glu S 290 Ala Leu Ser Ser Arg Ala Pro Ser Lys Pro His Val Ser Ser Pro Ser Leu Asn Leu Ala Gln Thr Ser Ser Phe Gly Pro Asp 310 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Ser Arg Ala Cys Glu S 350 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Arg Asn Pro Ile Pro Arg Arg Arg Arg Phe Leu Pro Arg Arg Arg Pro Ile Pro Pro Ile Pro Arg Pro Ile Pro Pro Pro Ile Pro Pro Pro Ile Pro Pro Pro Pro Ile Pro | Leu | Tyr L | | _ | Met | Pro | Gln | Pro | - | Ala | Ala | Gly | Asp | | Leu | Ser |
| 210 | Leu | | | Lys | Gly | Gln | Leu | | Ser | Ser | Pro | Ile | | Ser | Glu | Ser |
| 225 230 235 Asp Ser Ser Ser Arg Leu 245 Pro Arg His Trp Val Arg Pro 250 Arg Pro Val Ala Gen 250 Leu Ile Pro Asp Leu 260 Glu Val His Pro 265 Leu Glu Ile Tyr Arg Gen 270 Pro Arg Arg Ser Gln Ala Gly Thr Ala Thr Ser Ala Cys 285 Glu Ser 275 Ala Leu Ser Ser Arg Ala Pro 295 Ser Lys Pro His Val Ser Ser Ser Pro 300 Phe Pro Leu Gln Arg Cys Ala Thr Phe Arg Ala Leu Gly Pro Arg Ala Leu Gly Pro Arg 315 Ser Leu Asn Leu Ala Gln Thr Ser Pro Ser Phe Gly Ser Asn Value Ser Pro 340 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile Pro 340 | | | Sly | Ala | Leu | Thr | | Ala | Glu | Pro | Ser | | Phe | Gln | Glu | Pro |
| Leu Ile Pro Asp Leu Glu Val His Pro Leu Glu Ile Tyr Arg G 270 Pro Arg Arg Ser Gln Ala Gly Thr Ala Thr Ser Ala Cys Glu S 285 Ala Leu Ser Ser Arg Ala Pro Ser Lys Pro His Val Ser Ser Pro Ser Leu Asn Leu Ala Gln Thr Ser Pro Ser Pro Ser Phe Gly Ser Asn V 335 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile Pro Add Ser Ser Pro Ser Pro Arg Asn Pro Ile Pro Add Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Ser Pro Ser Pro Ser Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser Pro Ser Pro Arg Asn Pro Ile Pro Arg Ser Pro Ser P | | Val L | Leu | Gly | Glu | _ | Pro | Lys | His | Lys | | Thr | Thr | Leu | Arg | Met 240 |
| Pro Arg Arg Ser Gln Ala Gly Thr Ala Thr Ser Ala Cys Glu Ser Ala Leu Ser Ser Arg Ala Pro Ser Lys Pro His Val Ser Ser Ser Ser Pro Ser Pro His Val Ser Ser Ser Pro Ser Pro Ser Pro Ser Pro Pro Ala Leu Gly Pro Pro Arg Ala Leu Pro Pro Pro Arg Pro Pro Pro Arg Arg Pro Pro Arg Arg Pro Pro Arg P | Asp | Ser S | Ser | Arg | | Pro | Arg | His | Trp | | Arg | Pro | Val | Ala | Glu 255 | Val |
| Ala Leu Ser Ser Arg Ala Pro Ser Lys Pro His Val Ser Ser Pro 290 Phe Pro Leu Gln Arg Cys Ala Thr Phe Arg Ala Leu Gly Pro Assorting 310 Ser Leu Asn Leu Ala Gln Thr Ser Pro Ser Phe Gly Ser Asn Value Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile Pro 340 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile Pro 340 | Leu | Ile P | | _ | Leu | Glu | Val | His | | Leu | Glu | Ile | Tyr | _ | Gly | Arg |
| Phe Pro Leu Gln Arg Cys Ala Thr Phe Arg Ala Leu Gly Pro A 305 Ser Leu Asn Leu Ala Gln Thr Ser Pro Ser Phe Gly Ser Asn V 325 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile P 340 | Pro | - | _ | Ser | Gln | Ala | Gly | | Ala | Thr | Ser | Ala | _ | Glu | Ser | Gln |
| 305 310 315 Ser Leu Asn Leu Ala Gln Thr Ser Pro Ser Phe Gly Ser Asn V 325 330 Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile P 340 340 345 | Ala | | Ser | Ser | Arg | Ala | | Ser | Lys | Pro | His | | Ser | Ser | Pro | Arg |
| Phe Leu Ser Pro Gly Phe Arg Phe Leu Pro Arg Asn Pro Ile Page 340 345 350 | | Pro L | Leu | Gln | Arg | _ | Ala | Thr | Phe | Arg | | Leu | Gly | Pro | Asp | Pro 320 |
| 340 345 350 | Ser | Leu A | Asn | Leu | | Gln | Thr | Ser | Pro | | Phe | Gly | Ser | Asn | Val 335 | Pro |
| Asp Val Ala Ser Thr Pro Thr Pro Lys Leu Trp Pro Leu Ala L | Phe | Leu S | | | Gly | Phe | Arg | Phe | | Pro | Arg | Asn | Pro | | Pro | Pro |
| 355 360 365 | Asp | | | Ser | Thr | Pro | Thr | | Lys | Leu | Trp | Pro | | Ala | Lys | Trp |

Asp Ala Leu Glu Val Pro His Gly Gln Glu Gly Ser His Met Leu Ala

Pro Ser Gly Trp Glu Arg Glu Ala Glu Gln Leu Gly Glu Leu Trp Ala Gly Arg Thr Arg Val Pro Pro Gln Gly Gln Glu Pro Val Glu Val Thr Pro Leu Glu Glu Asp Ser Gly Trp Pro Leu Ala Ala Pro Gln Val Leu Glu Ala Thr Ser Gln Val Leu Trp Lys Pro Met Val Ile Ser Glu Thr Met Lys Leu Val Pro Gly Val Ser Met Trp Asn Arg Gly Thr Gln Glu Leu Leu Asn Pro Ala Val Ile Arg Lys Glu Ala Glu Glu Gly Thr Pro Gln Ala Pro Glu Gln Gln Pro Ile Gln Thr Gly Val Ser Lys Pro <210> 422 <211> 300 <212> PRT <213> Mus musculus <400> 422 Met Gly Leu Val Leu Arg Lys Met Leu Ser Ser Gly Val Cys Thr Ser Asn Val Gln Leu Pro Gly Lys Val Ala Ile Val Thr Gly Ala Asn Thr Gly Ile Gly Lys Glu Thr Ala Lys Asp Leu Ala Gln Arg Gly Ala Arg Val Tyr Leu Ala Cys Arg Asp Val Asp Lys Gly Glu Leu Ala Ala Arg

Asp Leu Ala Asp Thr Lys Ser Ile Arg Ala Phe Ala Lys Asp Phe Leu 85 90 95

Glu Ile Gln Ala Val Thr Gly Asn Ser Gln Val Phe Val Arg Lys Leu

Ala Glu Glu Lys His Leu His Leu Leu Ile Asn Asn Ala Gly Val Met

| | | | 100 | | | | | 105 | | | | | 110 | |
|-----|-----|-----|-----|-----|-----|-----|------|-----|-----|-----|-----|-----|-----|----|
| Mot | Cvc | Dro | Тих | Sor | Lvc | Thr | 71.7 | Λen | Glv | Dho | Glu | Mot | uic | Tl |

Met Cys Pro Tyr Ser Lys Thr Ala Asp Gly Phe Glu Met His Ile Gly
115 120 125

Val Asn His Leu Gly His Phe Leu Leu Thr His Leu Leu Glu Lys 130 135 140

Leu Lys Glu Ser Ala Pro Ser Arg Ile Val Asn Leu Ser Ser Leu Gly
145 150 155 160

His His Leu Gly Arg Ile His Phe His Asn Leu Gln Gly Glu Lys Phe 165 170 175

Tyr Ser Ala Gly Leu Ala Tyr Cys His Ser Lys Leu Ala Asn Ile Leu 180 185 190

Phe Thr Lys Glu Leu Ala Lys Arg Leu Lys Gly Ser Gly Val Thr Thr 195 200 205

Tyr Ser Val His Pro Gly Thr Val His Ser Glu Leu Thr Arg Tyr Ser 210 215 220

Ser Ile Met Arg Trp Leu Trp Gln Leu Phe Phe Val Phe Ile Lys Thr 225 230 235 240

Pro Gln Glu Gly Ala Gln Thr Ser Leu Tyr Cys Ala Leu Thr Glu Gly 245 250 255

Leu Glu Ser Leu Ser Gly Ser His Phe Ser Asp Cys Gln Leu Ala Trp
260 265 270

Val Ser Tyr Gln Gly Arg Asn Glu Ile Ile Ala Arg Arg Leu Trp Asp 275 280 285

Val Ser Cys Asp Leu Leu Gly Leu Pro Val Asp Trp 290 295 300

<210> 423

<211> 293

<212> PRT

<213> Mus musculus

<400> 423

Met Leu Ser Ser Gly Val Cys Thr Ser Asn Val Gln Leu Pro Gly Lys
1 5 10 15

| Val | Ala | Ile | Val 20 | Thr | Gly | Ala | Asn | Thr 25 | Gly | Ile | Gly | Lys | Glu 30 | Thr | Ala |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|-----------------|------------|
| Lys | Asp | Leu 35 | Ala | Gln | Arg | Gly | Ala 40 | Arg | Val | Tyr | Leu | Ala 45 | Cys | Arg | Asp |
| Val | Asp 50 | Lys | Gly | Glu | Leu | Ala 55 | Ala | Arg | Glu | Ile | Gln 60 | Ala | Val | Thr | Gly |
| Asn 65 | Ser | Gln | Val | Phe | Val 70 | Arg | Lys | Leu | Asp | Leu 75 | Ala | Asp | Thr | Lys | Ser 80 |
| Ile | Arg | Ala | Phe | Ala 85 | Lys | Asp | Phe | Leu | Ala 90 | Glu | Glu | Lys | His | Leu 95 | His |
| Leu | Leu | Ile | Asn 100 | Asn | Ala | Gly | Val | Met 105 | Met | Cys | Pro | Tyr | Ser 110 | Lys | Thr |
| Ala | Asp | Gly 115 | Phe | Glu | Met | His | Ile 120 | Gly | Val | Asn | His | Leu 125 | Gly | His | Phe |
| Leu | Leu 130 | Thr | His | Leu | Leu | Leu 135 | Glu | Lys | Leu | Lys | Glu 140 | Ser | Ala | Pro | Ser |
| Arg 145 | Ile | Val | Asn | Leu | Ser 150 | Ser | Leu | Gly | His | His 155 | Leu | Gly | Arg | Ile | His 160 |
| Phe | His | Asn | Leu | Gln 165 | Gly | Glu | Lys | Phe | Tyr 170 | Ser | Ala | Gly | Leu | Ala 175 | Tyr |
| Cys | His | Ser | Lys 180 | Leu | Ala | Asn | Ile | Leu 185 | Phe | Thr | Lys | Glu | Leu 190 | Ala | Lys |
| Arg | Leu | Lys 195 | Gly | Ser | Gly | Val | Thr 200 | Thr | Tyr | Ser | Val | His 205 | Pro | Gly | Thr |
| Val | His 210 | Ser | Glu | Leu | Thr | Arg 215 | Tyr | Ser | Ser | Ile | Met 220 | Arg | Trp | Leu | Trp |
| Gln 225 | Leu | Phe | Phe | Val | Phe 230 | Ile | Lys | Thr | Pro | Gln 235 | Glu | Gly | Ala | Gln | Thr 240 |
| Ser | Leu | Tyr | Cys | Ala 245 | Leu | Thr | Glu | Gly | Leu 250 | Glu | Ser | Leu | Ser | Gly 255 · | Ser |
| His | Phe | Ser | Asp 260 | Cys | Gln | Leu | Ala | Trp 265 | Val | Ser | Tyr | Gln | Gly 270 | Arg | Asn |

Glu Ile Ile Ala Arg Arg Leu Trp Asp Val Ser Cys Asp Leu Leu Gly 275 280 285

Leu Pro Val Asp Trp 290

<210> 424

<211> 316

<212> PRT

<213> Mus musculus

<400> 424

Met Phe Gly Phe Leu Leu Leu Ser Leu Pro Phe Ile Leu Tyr Leu
1 5 10 15

Val Thr Pro Lys Ile Arg Lys Met Leu Ser Ser Gly Val Cys Thr Ser 20 25 30

Asn Val Gln Leu Pro Gly Lys Val Ala Ile Val Thr Gly Ala Asn Thr 35 40 45

Gly Ile Gly Lys Glu Thr Ala Lys Asp Leu Ala Gln Arg Gly Ala Arg 50 55 60

Val Tyr Leu Ala Cys Arg Asp Val Asp Lys Gly Glu Leu Ala Ala Arg
65 70 75 80

Glu Ile Gln Ala Val Thr Gly Asn Ser Gln Val Phe Val Arg Lys Leu 85 90 95

Asp Leu Ala Asp Thr Lys Ser Ile Arg Ala Phe Ala Lys Asp Phe Leu
100 105 110

Ala Glu Glu Lys His Leu His Leu Leu Ile Asn Asn Ala Gly Val Met
115 120 125

Met Cys Pro Tyr Ser Lys Thr Ala Asp Gly Phe Glu Met His Ile Gly 130 135 140

Val Asn His Leu Gly His Phe Leu Leu Thr His Leu Leu Leu Glu Lys 145 150 155 160

Leu Lys Glu Ser Ala Pro Ser Arg Ile Val Asn Leu Ser Ser Leu Gly
165 170 175

His His Leu Gly Arg Ile His Phe His Asn Leu Gln Gly Glu Lys Phe 180 185 190 Tyr Ser Ala Gly Leu Ala Tyr Cys His Ser Lys Leu Ala Asn Ile Leu 195 200 Ship Ser Lys Gly Ser Gly Val Thr Thr 210 215 Ship Ser Ser Gly Val Thr Thr

Tyr Ser Val His Pro Gly Thr Val His Ser Glu Leu Thr Arg Tyr Ser 225 230 235 240

Ser Ile Met Arg Trp Leu Trp Gln Leu Phe Phe Val Phe Ile Lys Thr 245 250 255

Pro Gln Glu Gly Ala Gln Thr Ser Leu Tyr Cys Ala Leu Thr Glu Gly 260 265 270

Leu Glu Ser Leu Ser Gly Ser His Phe Ser Asp Cys Gln Leu Ala Trp 275 280 285

Val Ser Tyr Gln Gly Arg Asn Glu Ile Ile Ala Arg Arg Leu Trp Asp 290 295 300

Val Ser Cys Asp Leu Leu Gly Leu Pro Val Asp Trp 305 310 315

<210> 425

<211> 353

<212> PRT

<213> Mus musculus

<400> 425

Met Phe Gly Phe Leu Leu Leu Ser Leu Pro Phe Ile Leu Tyr Leu
1 5 10 15

Val Thr Pro Lys Ile Arg Lys Met Leu Ser Ser Gly Val Cys Thr Ser 20 25 30

Asn Val Gln Leu Pro Gly Lys Val Ala Ile Val Thr Gly Ala Asn Thr 35 40 45

Gly Ile Gly Lys Glu Thr Ala Lys Asp Leu Ala Gln Arg Gly Ala Arg
50 55 60

Val Tyr Leu Ala Cys Arg Asp Val Asp Lys Gly Glu Leu Ala Ala Arg 65 70 75 80

Glu Ile Gln Ala Val Thr Gly Asn Ser Gln Val Phe Val Arg Lys Leu

| Asp | Leu | Ala | Asp 100 | Thr | Lys | Ser | Ile | Arg 105 | Ala | Phe | Ala | Lys | Asp 110 | Phe | Leu |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ala | Glu | Glu 115 | Lys | His | Leu | His | Leu 120 | Leu | Ile | Asn | Asn | Ala 125 | Gly | Val | Met |
| Met | Cys 130 | Pro | Tyr | Ser | Lys | Thr 135 | Ala | Asp | Gly | Phe | Glu 140 | Met | His | Ile | Gly |
| Val 145 | Asn | His | Leu | Gly | His 150 | Phe | Leu | Leu | Thr | His 155 | Leu | Leu | Leu | Glu | Lys 160 |
| Leu | Lys | Glu | Ser | Ala 165 | Pro | Ser | Arg | Ile | Val 170 | Asn | Leu | Ser | Ser | Leu 175 | Gly |
| His | His | Leu | Gly 180 | Arg | Ile | His | Phe | His 185 | Asn | Leu | Gln | Gly | Glu 190 | Lys | Phe |
| Tyr | Ser | Ala 195 | Gly | Leu | Ala | Tyr | Cys 200 | His | Ser | Lys | Leu | Ala 205 | Asn | Ile | Leu |
| Phe | Thr 210 | Lys | Glu | Leu | Ala | Lys 215 | Arg | Leu | Lys | Gly | Ser 220 | Gly | Val | Thr | Thr |
| Tyr 225 | Ser | Val | His | Pro | Gly 230 | Thr | Val | His | Ser | Glu 235 | Leu | Thr | Gly | Tyr | Ser 240 |
| Ser | Ile | Met | Arg | Trp 245 | Leu | Trp | Gln | Leu | Phe 250 | Phe | Val | Phe | Ile | Lys 255 | Thr |
| Pro | Gln | Glu | Gly 260 | Ala | Gln | Thr | Ser | Leu 265 | Tyr | Cys | Ala | Leu | Thr 270 | Glu | Gly |
| Leu | Glu | Ser 275 | Leu | Ser | Gly | Arg | His 280 | Phe | Ser | Asp | Cys | Gln 285 | Leu | Ala | Trp |
| Val | Ser 290 | Tyr | Gln | Gly | Arg | Asn 295 | Glu | Ile | Ile | Ala | Arg 300 | Arg | Leu | Trp | Asp |
| Val 305 | Ser | Cys | Asp | Leu | Leu 310 | Ala | Ser | Gln | Trp | Ile 315 | Gly | Lys | Trp | Trp | Phe 320 |
| Gly | Pro | Lys | Arg | Arg 325 | Leu | Glu | Glu | Met | Met 330 | Ile | Ile | Leu | Gln | Ser 335 | Gly |
| Gln | Asn | Leu | Glu | Pro | Glu | Glu | Arg | Arg | Thr | Ser | Ser | Leu | Ser | Cys | Leu |

340 345 350

Ala

<210> 426

<211> 127

<212> PRT

<213> Homo sapiens

<400> 426

Thr Gly Lys Ile Ala Ile Val Thr Gly Ala Asn Ser Gly Ile Gly Lys
1 5 10 15

Val Val Ser Gln Asp Leu Ala Arg Cys Gly Ala Gln Val Ile Leu Thr 20 25 30

Cys Gln Ser Arg Glu Cys Gly Gln Gln Ala Leu Ala Glu Ile Gln Ala 35 40 45

Ala Ser Asn Ser Asn Arg Leu Leu Gly Glu Val Asp Leu Ser Ser 50 55 60

Met Thr Ser Ile Arg Ser Phe Ala Arg Arg Leu Leu Gln Glu Asn Pro 65 70 75 80

Glu Ile His Leu Leu Val Asn Asn Ala Gly Val Ser Gly Phe Arg Arg 85 90 95

His Leu Pro Gln Gly Ala Trp Ile Ser Pro Leu Ser Leu Thr Met Leu 100 105 110

Gly Pro Phe Cys Ser Gln Ile Tyr Ser Lys Asp Leu Lys Gln Gly
115 120 125

<210> 427

<211> 128

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: short chain
 dehydrogenase domain sequence

<400> 427

Thr Gly Lys Val Ala Leu Val Thr Gly Ala Ser Ser Gly Ile Gly Leu

Ala Ile Ala Lys Arg Leu Ala Glu Glu Gly Ala Lys Val Val Val 20 25 30

Asp Arg Arg Glu Glu Lys Ala Glu Ala Ala Glu Leu Lys Ala Glu
35 40 45

Leu Gly Asp Arg Ala Leu Phe Ile Gln Leu Asp Val Thr Asp Glu Glu 50 55 60

Ser Ile Lys Ala Ala Val Ala Gln Ala Val Glu Glu Leu Gly Arg Leu 65 70 75 80

Asp Val Leu Val Asn Asn Ala Gly Ile Leu Gly Pro Gly Glu Pro Phe 85 90 95

Glu Leu Ser Glu Asp Asp Trp Glu Arg Val Ile Asp Val Asn Leu Thr
100 105 110

Gly Val Phe Leu Leu Thr Gln Ala Val Leu Pro His Met Leu Lys Arg 115 120 125

<210> 428

<211> 158

<212> PRT

<213> Homo sapiens

<400> 428

Met Glu Val Met Asp Val Phe Ser Thr Asp Asp Leu Thr Gly Phe Leu

1 5 10 15

Gln Thr Lys Ala Gln Gln Gly Trp Leu Val Ala Gly Thr Val Gly Cys
20 25 30

Pro Ser Thr Glu Asp Pro Gln Ser Ser Glu Ile Pro Ile Met Ser Cys 35 40 45

Leu Glu Phe Leu Trp Glu Arg Pro Thr Leu Leu Val Leu Gly Asn Glu 50 55 60

Gly Ser Gly Leu Ser Gln Glu Val Gln Ala Ser Cys Gln Leu Leu 65 70 75 80

Thr Ile Leu Pro Arg Arg Gln Leu Pro Pro Gly Leu Glu Ser Leu Asn Val Ser Val Ala Ala Gly Ile Leu Leu His Ser Ile Cys Ser Gln Arg Lys Gly Phe Pro Thr Glu Gly Glu Arg Arg Gln Leu Leu Gln Asp Pro Gln Glu Pro Ser Ala Arg Ser Glu Gly Leu Ser Met Ala Gln His Pro Gly Leu Ser Ser Gly Pro Glu Lys Glu Arg Gln Asn Glu Gly <210> 429 <211> 155 <212> PRT <213> Homo sapiens <400> 429 Met Asp Val Phe Ser Thr Asp Asp Leu Thr Gly Phe Leu Gln Thr Lys Ala Gln Gln Gly Trp Leu Val Ala Gly Thr Val Gly Cys Pro Ser Thr Glu Asp Pro Gln Ser Ser Glu Ile Pro Ile Met Ser Cys Leu Glu Phe Leu Trp Glu Arg Pro Thr Leu Leu Val Leu Gly Asn Glu Gly Ser Gly Leu Ser Gln Glu Val Gln Ala Ser Cys Gln Leu Leu Leu Thr Ile Leu Pro Arg Arg Gln Leu Pro Pro Gly Leu Glu Ser Leu Asn Val Ser Val Ala Ala Gly Ile Leu Leu His Ser Ile Cys Ser Gln Arg Lys Gly Phe Pro Thr Glu Gly Glu Arg Arg Gln Leu Leu Gln Asp Pro Gln Glu Pro Ser Ala Arg Ser Glu Gly Leu Ser Met Ala Gln His Pro Gly Leu Ser

<210> 430 <211> 124 <212> PRT <213> Homo sapiens <400> 430 Met Asp Val Phe Ala Thr Pro Asp Leu Pro Gly Phe Leu Gln Ala Lys Ala Gln Gln Gly Trp Leu Val Val Gly Thr Val Gly Cys Pro Gly Pro Glu Ile Ser Gln Ser Ser Lys Val Pro Ile Thr Ser Cys Leu Glu Phe Val Trp Asp Arg Pro Thr Leu Leu Val Leu Gly Ser Glu Gly Ser Gly Leu Ser Gln Glu Val Phe Ala Ser Cys Gln Leu Leu Thr Ile Leu Pro Arg Arg His Leu Pro Pro Gly Leu Glu Ser Leu Asn Val Ser Val Ala Thr Gly Ile Leu Leu His Ser Ile Cys Ser Gln Lys Lys Gly Phe Pro Val Gln Glu Arg Gly Gln Leu Leu Gln Asp Ser <210> 431 <211> 181 <212> PRT <213> Homo sapiens <400> 431 Met Phe Ser Ala Ile Arg Ser Gln His Ser Gly Val Asp Ile Cys Ile Asn Asn Ala Gly Leu Ala Arg Pro Asp Thr Leu Leu Ser Gly Ser Thr

Ser Gly Pro Glu Lys Glu Arq Gln Asn Glu Gly

Cys Thr Arg Glu Ala Tyr Gln Ser Met Lys Glu Arg Asn Val Asp Asp Gly His Ile Ile Asn Ile Asn Ser Met Ser Gly His Arg Val Leu Pro Leu Ser Val Thr His Phe Tyr Ser Ala Thr Lys Tyr Ala Val Thr Ala Leu Thr Glu Gly Leu Arg Gln Glu Leu Arg Glu Ala Gln Thr His Ile Arg Ala Thr Cys Ile Ser Pro Gly Val Val Glu Thr Gln Phe Ala Phe Lys Leu His Asp Lys Asp Pro Glu Lys Ala Ala Ala Thr Tyr Glu Gln Met Lys Cys Leu Lys Pro Glu Asp Val Ala Glu Ala Val Ile Tyr Val Leu Ser Thr Pro Ala His Ile Gln Ile Gly Asp Ile Gln Met Arg Pro Thr Glu Gln Val Thr <210> 432 <211> 181 <212> PRT <213> Mus musculus <400> 432 Met Ser Ser Ala Ile Arg Ser Gln His Ser Gly Val Asp Ile Cys Ile Asn Asn Ala Gly Leu Ala Arg Pro Asp Thr Leu Leu Ser Gly Ser Thr Ser Gly Trp Lys Asp Met Phe Asn Val Asn Val Leu Ala Leu Ser Ile Cys Thr Arg Glu Ala Tyr Gln Ser Met Lys Glu Arg Asn Val Asp Asp

Ser Gly Trp Lys Asp Met Phe Asn Val Asn Val Leu Ala Leu Ser Ile

Gly His Ile Ile Asn Ile Asn Ser Met Ser Gly His Arg Val Leu Pro 65 70 75 80

Leu Ser Val Thr His Phe Tyr Ser Ala Thr Lys Tyr Ala Val Thr Ala 85 90 95

Leu Thr Glu Gly Leu Arg Gln Glu Leu Arg Glu Ala Gln Thr His Ile 100 105 110

Arg Ala Thr Cys Ile Ser Pro Gly Val Val Glu Thr Gln Phe Ala Phe 115 120 125

Lys Leu His Asp Lys Asp Pro Glu Lys Ala Ala Ala Thr Tyr Glu Gln 130 135 140

Leu Ser Thr Pro Ala His Ile Gln Ile Gly Asp Ile Gln Met Arg Pro 165 170 175

Thr Glu Gln Val Thr 180

<210> 433

<211> 182

<212> PRT

<213> Homo sapiens

<400> 433

Asp Arg Leu Ala Leu Val Thr Gly Ala Ser Gly Gly Ile Gly Ala Ala 1 5 10 15

Val Ala Arg Ala Leu Val Gln Gln Gly Leu Lys Val Val Gly Cys Ala 20 25 30

Arg Thr Val Gly Asn Ile Glu Glu Leu Ala Ala Glu Cys Lys Ser Ala 35 40 45

Gly Tyr Pro Gly Thr Leu Ile Pro Tyr Arg Cys Asp Leu Ser Asn Glu
50 55 60

Glu Asp Ile Leu Ser Met Phe Ser Ala Ile Arg Ser Gln His Ser Gly
65 70 75 80

Val Asp Ile Cys Ile Asn Asn Ala Gly Leu Ala Arg Pro Asp Thr Leu

85 90 95

Leu Ser Gly Ser Thr Ser Gly Trp Lys Asp Met Phe Asn Val Asn Val
100 105 110

Leu Ala Leu Ser Ile Cys Thr Arg Glu Ala Tyr Gln Ser Met Lys Glu
115 120 125

Arg Asn Val Asp Asp Gly His Ile Ile Asn Ile Asn Ser Met Ser Gly 130 135 140

His Arg Val Leu Pro Leu Ser Val Thr His Phe Tyr Ser Ala Thr Lys 145 150 155 160

Tyr Ala Val Thr Ala Leu Thr Glu Gly Leu Arg Gln Glu Leu Arg Glu 165 170 175

Ala Gln Thr His Ile Arg 180

<210> 434

<211> 174

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: short chain dehydrogenase domain sequence

<400> 434

Gly Lys Val Ala Leu Val Thr Gly Ala Ser Ser Gly Ile Gly Leu Ala 1 5 10 15

Ile Ala Lys Arg Leu Ala Glu Glu Gly Ala Lys Val Val Val Asp 20 25 30

Arg Arg Glu Glu Lys Ala Glu Ala Ala Glu Leu Lys Ala Glu Leu
35 40 45

Gly Asp Arg Ala Leu Phe Ile Gln Leu Asp Val Thr Asp Glu Glu Ser 50 55 60

Ile Lys Ala Ala Val Ala Gln Ala Val Glu Glu Leu Gly Arg Leu Asp
65 70 75 80

Val Leu Val Asn Asn Ala Gly Ile Leu Gly Pro Gly Glu Pro Phe Glu 85 90 95 Leu Ser Glu Asp Asp Trp Glu Arg Val Ile Asp Val Asn Leu Thr Gly 100 105 110

Val Phe Leu Leu Thr Gln Ala Val Leu Pro His Met Leu Lys Arg Ser 115 120 125

Gly Gly Arg Ile Val Asn Ile Ser Ser Val Ala Gly Leu Val Pro Ser 130 135 140

Pro Gly Leu Ser Ala Tyr Ser Ala Ser Lys Ala Ala Val Val Gly Phe 145 150 155 160

Thr Arg Ser Leu Ala Leu Glu Leu Ala Pro His Gly Ile Arg 165 170

<210> 435

<211> 115

<212> PRT

<213> Homo sapiens

<400> 435

Leu Val Leu Asp Gly Ile Gln Asp Pro Arg Asn Phe Gly Ala Val Leu 1 5 10 15

Arg Ser Ala His Phe Leu Gly Val Asp Lys Thr Lys Ala Gln Gln Gly 20 25 30

Trp Leu Val Ala Gly Thr Val Gly Cys Pro Ser Thr Glu Asp Pro Gln 35 40 45

Ser Ser Glu Ile Pro Ile Met Ser Cys Leu Glu Phe Leu Trp Glu Arg
50 55 60

Pro Thr Leu Leu Val Leu Gly Asn Glu Gly Ser Gly Leu Ser Gln Glu 65 70 75 80

Val Gln Ala Ser Cys Gln Leu Leu Thr Ile Leu Pro Arg Arg Gln 85 90 95

Leu Pro Pro Gly Leu Glu Ser Leu Asn Val Ser Val Ala Ala Gly Ile 100 105 110

Leu Leu His

```
<210> 436
<211> 140
<212> PRT
<213> Artificial Sequence
<220>
<223> Description of Artificial Sequence: SpoU rRNA
      Methylase family domain sequence
<400> 436
Val Val Leu Asp Glu Val Glu Ile Pro His Asn Ile Gly Ala Ile Ile
                                      10
Arg Thr Cys Ala Ala Leu Gly Val Asp Gly Ile Val Ile Val Asp Asp
             20
                                  25
Gly Phe Ala Leu Leu Asp Arg Arg Leu Arg Arg Ala Ser Leu Gly Tyr
                             40
Ala Glu Ser Val Pro Val Ile Arg Val Asp Asn Leu Glu Glu Phe Leu
                         55
                                              60
Ala His Leu Lys Glu Ser Gly Ile Trp Leu Leu Thr Thr Ser Gly Asp
 65
                                          75
                     70
Gly Asn Ala Asp Pro Leu Asp Tyr Glu Asp Gly Ala Lys Arg Leu Ala
                 85
                                      90
Leu Val Phe Gly Ser Glu Thr Thr Gly Leu Ser Asn Leu Ala Leu Glu
            100
                                 105
                                                     110
Pro Ala Asp Gln Arg Ile Arg Ile Pro Met Asn Gly Asp Val Arg Ser
        115
                            120
                                                 125
Leu Asn Val Ser Val Ala Val Gly Leu Leu Leu Tyr
    130
                        135
                                             140
<210> 437
<211> 159
<212> PRT
<213> Homo sapiens
```

<400> 437

Leu Val Thr Gly Ala Ser Gly Gly Ile Gly Ala Ala Val Ala Arg Ala 1 5 10 15

Leu Val Gln Gln Gly Leu Lys Val Val Gly Cys Ala Arg Thr Val Gly

20 25 30

Asn Ile Glu Glu Leu Ala Ala Glu Cys Lys Ser Ala Gly Tyr Pro Gly
35 40 45

Thr Leu Ile Pro Tyr Arg Cys Asp Leu Ser Asn Glu Glu Asp Ile Leu 50 55 60

Ser Met Phe Ser Ala Ile Arg Ser Gln His Ser Gly Val Asp Ile Cys 65 70 75 80

Ile Asn Asn Ala Gly Leu Ala Arg Pro Asp Thr Leu Leu Ser Gly Ser 85 90 95

Thr Ser Gly Trp Lys Asp Met Phe Asn Val Asn Val Leu Ala Leu Ser 100 105 110

Ile Cys Thr Arg Glu Ala Tyr Gln Ser Met Lys Glu Arg Asn Val Asp 115 120 125

Asp Gly His Ile Ile Asn Ile Asn Ser Met Ser Gly His Arg Val Leu 130 135 140

<210> 438

<211> 152

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: NAD dependent epimerase/dehydratase family domain sequence

<400> 438

Leu Val Thr Gly Gly Ala Gly Phe Ile Gly Ser His Leu Val Arg Glu
1 5 10 15

Leu Leu Asn Asn Gly Asp Asp Lys Val Val Leu Asp Asn Leu Thr
20 25 30

Tyr Ala Gly Asn Glu Ala Arg Leu Arg Val Ile Glu Gly Gly Pro Arg
35 40 45

Tyr Thr Phe Val Lys Gly Asp Ile Cys Asp Arg Asp Leu Leu Asp Lys 50 55 60

Val Phe Ala Glu Asn Gln Pro Asp Ala Val Ile His Phe Ala Ala Glu 70 75 65 Ser His Val Asp Arg Ser Ile Glu Lys Pro Leu Ala Tyr Ile Asp Thr 90 85 Asn Val Val Gly Thr Leu Thr Leu Leu Glu Ala Ala Arg Lys Ala Gly 100 105 110 Val Phe Lys Phe Val Phe Ser Ser Thr Asp Glu Val Tyr Gly Asp Leu 120 Pro Ser Ile Pro Ile Thr Glu Asp Thr Pro Tyr Gly Pro Ser Ser Pro 135 140 Tyr Gly Ala Ser Lys Ala Ser Ser 150 <210> 439 <211> 796 <212> PRT <213> Homo sapiens <400> 439 Met Glu Ala Gly Gly Glu Arg Phe Leu Arg Gln Arg Gln Val Leu Leu 5 10 15 Leu Phe Val Phe Leu Gly Gly Ser Leu Ala Gly Ser Glu Ser Arg Arg 20 25 30 Tyr Ser Val Ala Glu Glu Lys Glu Lys Gly Phe Leu Ile Ala Asn Leu 35 40 Ala Lys Asp Leu Gly Leu Arg Val Glu Glu Leu Ala Ala Arg Gly Ala 50 55 60 Gln Val Val Ser Lys Gly Asn Lys Gln His Phe Gln Leu Ser His Gln 70 75 Thr Gly Asp Leu Leu Asn Glu Lys Leu Asp Arg Glu Glu Leu Cys

Pro Leu Gln Phe Val Thr Asn Glu Leu Arg Ile Ile Asp Val Asn Asp

Gly Pro Thr Glu Pro Cys Ile Leu His Phe Gln Ile Leu Leu Gln Asn 100 105 110

| | | 115 | | | | | 120 | | | | | 125 | | | |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| His | Ser 130 | Pro | Val | Phe | Phe | Glu 135 | Asn | Glu | Met | His | Leu 140 | Lys | Ile | Leu | Glu |
| Ser 145 | Thr | Leu | Pro | Gly | Thr 150 | Val | Ile | Pro | Leu | Gly 155 | Asn | Ala | Glu | Asp | Leu 160 |
| Asp | Val | Gly | Arg | Asn 165 | Ser | Leu | Gln | Asn | Tyr 170 | Thr | Ile | Thr | Pro | Asn 175 | Ser |
| His | Phe | His | Val 180 | Leu | Thr | Arg | Ser | Arg 185 | Arg | Asp | Gly | Arg | Lys 190 | Tyr | Pro |
| Glu | Leu | Val 195 | Leu | Asp | Lys | Ala | Leu 200 | Asp | Arg | Glu | Glu | Gln 205 | Pro | Glu | Leu |
| Ser | Leu 210 | Thr | Leu | Thr | Ala | Leu 215 | Asp | Gly | Gly | Ser | Pro 220 | Pro | Arg | Ser | Gly |

Thr Ala Gln Ile Asn Ile Gln Val Leu Asp Ile Asn Asp Asn Ala Pro 225 230 235 240

Glu Phe Ala Gln Pro Leu Tyr Glu Val Ala Val Leu Glu Asn Thr Pro 245 250 255

Val Asn Ser Val Ile Val Thr Val Ser Ala Ser Asp Leu Asp Thr Gly 260 265 270

Ser Phe Gly Thr Ile Ser Tyr Ala Phe Phe His Ala Ser Glu Glu Ile 275 280 285

Arg Lys Thr Phe Gln Leu Asn Pro Ile Thr Gly Asp Met Gln Leu Val 290 295 300

Lys Tyr Leu Asn Phe Glu Ala Ile Asn Ser Tyr Glu Val Asp Ile Glu 305 310 315 320

Ala Lys Asp Gly Gly Leu Ser Gly Lys Ser Thr Val Ile Val Gln 325 330 335

Val Val Asp Val Asn Asp Asn Pro Pro Glu Leu Thr Leu Ser Ser Val 340 345 350

Asn Ser Pro Ile Pro Glu Asn Ser Gly Glu Thr Val Leu Ala Val Phe 355 360 365

Ser Val Ser Asp Leu Asp Ser Gly Asp Asn Gly Arg Val Met Cys Ser

| 370 | 375 | 380 |
|-----|-----|-----|
| | | |

| Ile 385 | Glu | Asn | Asn | Leu | Pro 390 | Phe | Phe | Leu | Lys | Pro 395 | Ser | Val | Glu | Asn | Phe 400 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Tyr | Thr | Leu | Val | Ser 405 | Glu | Gly | Ala | Leu | Asp 410 | Arg | Glu | Thr | Arg | Ser 415 | Glu |
| Tyr | Asn | Ile | Thr 420 | Ile | Thr | Ile | Thr | Asp 425 | Leu | Gly | Thr | Pro | Arg 430 | Leu | Lys |
| Thr | Lys | Tyr 435 | Asn | Ile | Thr | Val | Leu 440 | Val | Ser | Asp | Val | Asn 445 | Asp | Asn | Ala |
| Pro | Ala 450 | Phe | Thr | Gln | Ile | Ser 455 | Tyr | Thr | Leu | Phe | Val 460 | Arg | Glu | Asn | Asn |
| Ser 465 | Pro | Ala | Leu | His | Ile 470 | Gly | Ser | Val | Ser | Ala 475 | Thr | Asp | Arg | Asp | Ser 480 |
| Gly | Thr | Asn | Ala | Gln 485 | Val | Thr | Tyr | Ser | Leu 490 | Leu | Pro | Pro | Gln | Asp 495 | Pro |
| His | Leu | Pro | Leu 500 | Ser | Ser | Leu | Val | Ser 505 | Ile | Asn | Ala | Asp | Asn 510 | Gly | His |
| Leu | Phe | Ala 515 | Leu | Arg | Ser | Leu | Asp 520 | Tyr | Glu | Ala | Leu | Gln 525 | Ala | Phe | Glu |
| Phe | Arg 530 | Val | Gly | Ala | Thr | Asp 535 | Arg | Gly | Ser | Pro | Ala 540 | Leu | Ser | Ser | Glu |
| Ala 545 | Leu | Val | Arg | Val | Leu 550 | Val | Leu | Asp | Ala | Asn 555 | Asp | Asn | Ser | Pro | Phe 560 |
| Val | Leu | Tyr | Pro | Leu 565 | Gln | Asn | Gly | Ser | Ala 570 | Pro | Cys | Thr | Glu | Leu 575 | Val |
| Pro | Arg | Ala | Ala 580 | Glu | Pro | Gly | Tyr | Leu 585 | Val | Thr | Lys | Val | Val 590 | Ala | Val |
| Asp | Gly | Asp 595 | Ser | Gly | Gln | Asn | Ala 600 | Trp | Leu | Ser | Tyr | Gln 605 | Leu | Leu | Lys |
| Ala | Thr 610 | Glu | Pro | Gly | Leu | Phe 615 | Gly | Val | Trp | Ala | His 620 | Asn | Gly | Glu | Val |
| Arg | Thr | Ala | Arg | Leu | Leu | Ser | Glu | Arg | Asp | Ala | Ala | Lys | His | Arg | Leu |

Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg Ser Ala Thr Ala 645 650 655

Thr Leu His Val Leu Leu Val Asp Gly Phe Ser Gln Pro Tyr Leu Pro 660 665 670

Leu Pro Glu Ala Ala Pro Ala Gln Ala Gln Ala Asp Leu Leu Thr Val 675 680 685

Tyr Leu Val Val Ala Leu Ala Ser Val Ser Ser Leu Phe Leu Phe Ser 690 695 700

Val Leu Leu Phe Val Ala Val Arg Leu Cys Arg Arg Ser Arg Ala Ala 705 710 715 720

Ser Val Gly Arg Cys Ser Val Pro Glu Gly Pro Phe Pro Gly Gln Met 725 730 735

Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu
740 745 750

Val Cys Leu Thr Gly Gly Ser Gly Thr Asn Glu Phe Lys Phe Leu Lys
755 760 765

Pro Ile Ile Pro Asn Phe Val Ala Gln Gly Ala Glu Arg Val Ser Glu 770 780

Ala Asn Pro Ser Phe Arg Lys Ser Phe Glu Phe Ser 785 790 795

<210> 440

<211> 798

<212> PRT

<213> Homo sapiens

<400> 440

Met Glu Ala Gly Glu Gly Lys Glu Arg Val Pro Lys Gln Arg Gln Val
1 5 10 15

Leu Ile Phe Phe Val Leu Leu Gly Ile Ala Gln Ala Ser Cys Gln Pro 20 25 30

Arg His Tyr Ser Val Ala Glu Glu Thr Glu Ser Gly Ser Phe Val Ala 35 40 45

| Asn | Leu 50 | Leu | Lys | Asp | Leu | Gly 55 | Leu | Glu | Ile | Gly | Glu 60 | Leu | Ala | Val | Arg |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Gly 65 | Ala | Arg | Val | Val | Ser 70 | Lys | Gly | Lys | Lys | Met 75 | His | Leu | Gln | Phe | Asp 80 |
| Arg | Gln | Thr | Gly | Asp 85 | Leu | Leu | Leu | Asn | Glu 90 | Lys | Leu | Asp | Arg | Glu 95 | Glu |
| Leu | Cys | Gly | Pro 100 | Thr | Glu | Pro | Cys | Val 105 | Leu | Pro | Phe | Gln | Val 110 | Leu | Leu |
| Glu | Asn | Pro 115 | Leu | Gln | Phe | Phe | Gln 120 | Ala | Glu | Leu | Arg | Ile 125 | Arg | Asp | Val |
| Asn | Asp 130 | His | Ser | Pro | Val | Phe 135 | Leu | Asp | Lys | Glu | Ile 140 | Leu | Leu | Lys | Ile |
| Pro 145 | Glu | Ser | Ile | Thr | Pro 150 | Gly | Thr | Thr | Phe | Leu 155 | Ile | Glu | Arg | Ala | Gln 160 |
| Asp | Leu | Asp | Val | Gly 165 | Thr | Asn | Ser | Leu | Gln 170 | Asn | Tyr | Thr | Ile | Ser 175 | Pro |
| Asn | Phe | His | Phe 180 | His | Leu | Asn | Leu | Gln 185 | Asp | Ser | Leu | Asp | Gly 190 | Ile | Ile |
| Leu | Pro | Gln 195 | Leu | Val | Leu | Asn | Arg 200 | Ala | Leu | Asp | Arg | Glu 205 | Glu | Gln | Pro |
| Glu | Ile 210 | Arg | Leu | Thr | Leu | Thr 215 | Ala | Leu | Asp | Gly | Gly 220 | Ser | Pro | Pro | Arg |
| Ser 225 | Gly | Thr | Ala | Leu | Val 230 | Arg | Ile | Glu | Val | Val 235 | Asp | Ile | Asn | Asp | Asn 240 |
| Val | Pro | Glu | Phe | Ala 245 | Lys | Leu | Leu | Tyr | Glu 250 | Val | Gln | Ile | Pro | Glu 255 | Asp |
| Ser | Pro | Val | Gly 260 | Ser | Gln | Val | Ala | Ile 265 | | Ser | Ala | Arg | Asp 270 | Leu | Asp |
| Ile | Gly | Thr 275 | Asn | Gly | Glu | Ile | Ser 280 | Tyr | Ala | Phe | Ser | Gln 285 | Ala | Ser | Glu |
| Asp | Ile | Arg | Lys | Thr | Phe | Arg | Leu | Ser | Ala | Lys | Ser | Gly | Glu | Leu | Leu |

| Leu Arg 305 | Gln L | ys Le | 310 | Phe | Glu | Ser | Ile | Gln 315 | Thr | Tyr | Thr | Val | Asn 320 |
|----------------|-------|----------------------------|--------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ile Gln | Ala T | hr Ası 32 | _ | Gly | Gly | Leu | Ser 330 | Gly | Thr | Cys | Val | Val 335 | Phe |
| Val Gln | | let As _l 340 | D Leu | Asn | Asp | Asn 345 | Pro | Pro | Glu | Leu | Thr 350 | Met | Ser |
| Thr Leu | Ile A | sn Gl | n Ile | Pro | Glu 360 | Asn | Leu | Gln | Asp | Thr 365 | Leu | Ile | Ala |
| Val Phe 370 | Ser V | al Se | Asp | Pro 375 | Asp | Ser | Gly | Asp | Asn 380 | Gly | Arg | Met | Val |
| Cys Ser 385 | Ile G | iln As _l | 390 | Leu | Pro | Phe | Phe | Leu 395 | Lys | Pro | Ser | Val | Glu 400 |
| Asn Phe | Tyr T | hr Le | | Ile | Ser | Thr | Ala 410 | Leu | Asp | Arg | Glu | Thr 415 | Arg |
| Ser Glu | _ | sn Il | e Thr | Ile | Thr | Val 425 | Thr | Asp | Phe | Gly | Thr 430 | Pro | Arg |
| Leu Lys | Thr G | Slu Hi | s Asn | Ile | Thr 440 | Val | Leu | Val | Ser | Asp 445 | Val | Asn | Asp |
| Asn Ala 450 | Pro A | ala Ph | e Thr | Gln 455 | Thr | Ser | Tyr | Thr | Leu 460 | Phe | Val | Arg | Glu |
| Asn Asn 465 | Ser P | ro Al | 470 | His | Ile | Gly | Ser | Val 475 | Ser | Ala | Thr | Asp | Arg 480 |
| Asp Ser | Gly T | hr As: | | Gln | Val | Thr | Tyr 490 | Ser | Leu | Leu | Pro | Pro 495 | Gln |
| Asp Pro | | eu Pr | o Leu | Ala | Ser | Leu 505 | Val | Ser | Ile | Asn | Ala 510 | Asp | Asn |
| Gly His | Leu P | he Al | a Leu | Gln | Ser 520 | Leu | Asp | Tyr | Glu | Ala 525 | Leu | Gln | Ala |
| Phe Glu 530 | Phe A | irg Va | l Gly | Ala 535 | Ala | Asp | Arg | Gly | Ser 540 | Pro | Ala | Leu | Ser |
| Ser Glu 545 | Ala L | eu Va | L Arg 550 | Val | Leu | Val | Leu | Asp 555 | Ala | Asn | Asp | Asn | Ser 560 |

Pro Phe Val Leu Tyr Pro Leu Gln Asn Gly Ser Ala Pro Cys Thr Glu Leu Val Pro Arg Ala Ala Glu Pro Gly Tyr Leu Val Thr Lys Val Val Ala Val Asp Gly Asp Ser Gly Gln Asn Ala Trp Leu Ser Tyr Gln Leu Leu Lys Ala Thr Glu Pro Gly Leu Phe Gly Val Trp Ala His Asn Gly Glu Val Arg Thr Ala Arg Leu Leu Arg Glu Arg Asp Ala Ala Lys Gln Arg Leu Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg Ser Ala Thr Ala Thr Leu His Val Leu Leu Val Asp Gly Phe Ser Gln Pro Tyr Leu Leu Pro Glu Ala Ala Pro Ala Gln Ala Gln Ala Asp Leu Leu Thr Val Tyr Leu Val Val Ala Leu Ala Ser Val Ser Ser Leu Phe Leu Phe Ser Val Leu Leu Phe Val Ala Val Arg Leu Cys Arg Arg Ser Arg Ala Ala Ser Val Gly Arg Cys Ser Val Pro Glu Gly Pro Phe Pro Gly Gln Met Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu Val Cys Leu Thr Gly Gly Ser Gly Thr Asn Glu Phe Lys Phe Leu Lys Pro Ile Ile Pro Asn Phe Val Ala Gln Gly Ala Glu Arg Val Ser Glu Ala Asn Pro Ser Phe Arg Lys Ser Phe Glu Phe Thr

<210> 441 <211> 776

<212> PRT <213> Homo sapiens

<400> 441

Met Glu Ile Gly Trp Met His Asn Arg Arg Gln Arg Gln Val Leu Val
1 5 10 15

Phe Phe Val Leu Ser Leu Ser Gly Ala Gly Ala Glu Leu Gly Ser 20 25 30

Tyr Ser Val Val Glu Glu Thr Glu Arg Gly Ser Phe Val Ala Asn Leu
35 40 45

Gly Lys Asp Leu Gly Leu Gly Leu Thr Glu Met Ser Thr Arg Lys Ala
50 55 60

Arg Ile Ile Ser Gln Gly Asn Lys Gln His Leu Gln Leu Lys Ala Gln 65 70 75 80

Thr Gly Asp Leu Leu Ile Asn Glu Lys Leu Asp Arg Glu Glu Leu Cys
85 90 95

Gly Pro Thr Glu Pro Cys Ile Leu His Phe Gln Val Leu Met Glu Asn 100 105 110

Pro Leu Glu Ile Phe Gln Ala Glu Leu Arg Val Ile Asp Ile Asn Asp 115 120 125

His Ser Pro Met Phe Thr Glu Lys Glu Met Ile Leu Lys Ile Pro Glu 130 135 140

Asn Ser Pro Leu Gly Thr Glu Phe Pro Leu Asn His Ala Leu Asp Leu 145 150 155 160

Asp Val Gly Ser Asn Asn Val Gln Asn Tyr Lys Ile Ser Pro Ser Ser 165 170 175

His Phe Arg Val Leu Ile His Glu Phe Arg Asp Gly Arg Lys Tyr Pro 180 185 190

Glu Leu Val Leu Asp Lys Glu Leu Asp Arg Glu Glu Glu Pro Gln Leu 195 200 205

Arg Leu Thr Leu Thr Ala Leu Asp Gly Gly Ser Pro Pro Arg Ser Gly 210 215 220

Thr Ala Gln Val Arg Ile Glu Val Val Asp Ile Asn Asp Asn Ala Pro 225 230 235 240

| Glu | Phe | Glu | Gln | Pro 245 | Ile | Tyr | Lys | Val | Gln 250 | Ile | Pro | Glu | Asn | Ser 255 | Pro |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Leu | Gly | Ser | Leu 260 | Val | Ala | Thr | Val | Ser 265 | Ala | Arg | Asp | Leu | Asp 270 | Gly | Gly |
| Ala | Asn | Gly 275 | Lys | Ile | Ser | Tyr | Thr 280 | Leu | Phe | Gln | Pro | Ser 285 | Glu | Asp | Ile |
| Ser | Lys 290 | Thr | Leu | Glu | Val | Asn 295 | Pro | Met | Thr | Gly | Glu 300 | Val | Arg | Leu | Arg |
| Lys 305 | Gln | Val | Asp | Phe | Glu 310 | Met | Val | Thr | Ser | Tyr 315 | Glu | Val | Arg | Ile | Lys 320 |
| Ala | Thr | Asp | Gly | Gly 325 | Gly | Leu | Ser | Gly | Lys 330 | Cys | Thr | Leu | Leu | Leu 335 | Gln |
| Val | Val | Asp | Val 340 | Asn | Asp | Asn | Pro | Pro 345 | Gln | Val | Thr | Met | Ser 350 | Ala | Leu |
| Thr | Ser | Pro 355 | Ile | Pro | Glu | Asn | Ser 360 | Pro | Glu | Ile | Val | Val 365 | Ala | Val | Phe |
| Ser | Val 370 | Ser | Asp | Pro | Asp | Ser 375 | Gly | Asn | Asn | Gly | Lys 380 | Thr | Ile | Ser | Ser |
| Ile 385 | Gln | Glu | Asp | Leu | Pro 390 | Phe | Leu | Leu | Lys | Pro 395 | Ser | Val | Lys | Asn | Phe 400 |
| Tyr | Thr | Leu | Val | Thr 405 | Glu | Arg | Ala | Leu | Asp 410 | Arg | Glu | Ala | Arg | Ala 415 | Glu |
| Tyr | Asn | Ile | Thr 420 | Leu | Thr | Val | Thr | Asp 425 | Met | Gly | Thr | Pro | Arg 430 | Leu | Lys |
| Thr | Glu | His 435 | Asn | Ile | Thr | Val | Gln 440 | Ile | Ser | Asp | Val | Asn 445 | Asp | Asn | Ala |
| Pro | Thr 450 | Phe | Thr | Gln | Thr | Ser 455 | Tyr | Thr | Leu | Phe | Val 460 | Arg | Glu | Asn | Asn |
| Ser 465 | Pro | Ala | Leu | His | Ile 470 | Gly | Ser | Val | Ser | Ala 475 | Thr | Asp | Arg | Asp | Ser 480 |
| Gly | Thr | Asn | Ala | Gln 485 | Val | Thr | Tyr | Ser | Leu 490 | Leu | Pro | Pro | Gln | Asp 495 | Pro |

His Leu Pro Leu Ala Ser Leu Val Ser Ile Asn Ala Asp Asn Gly His Leu Phe Ala Leu Arg Ser Leu Asp Tyr Glu Ala Leu Gln Ala Phe Glu Phe Arg Val Gly Ala Thr Asp Arg Gly Ser Pro Ala Leu Ser Arg Glu Ala Leu Val Arg Val Leu Val Leu Asp Ala Asn Asp Asn Ser Pro Phe Val Leu Tyr Pro Leu Gln Asn Gly Ser Ala Pro Cys Thr Glu Leu Val Pro Arg Ala Ala Glu Pro Gly Tyr Leu Val Thr Lys Val Val Ala Val Asp Gly Asp Ser Gly Gln Asn Ala Trp Leu Ser Tyr Gln Leu Leu Lys Ala Thr Glu Pro Gly Leu Phe Gly Val Trp Ala His Asn Gly Glu Val Arg Thr Ala Arg Leu Leu Ser Glu Arg Asp Ala Ala Lys Gln Arg Leu Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg Ser Ala Thr Ala Thr Leu His Val Leu Leu Val Asp Gly Phe Ser Gln Pro Phe Leu Pro Leu Pro Glu Ala Ala Pro Gly Gln Thr Gln Ala Asn Ser Leu Thr Val Tyr Leu Val Val Ala Leu Ala Ser Val Ser Ser Leu Phe Leu Phe Ser Val Leu Leu Phe Val Ala Val Arg Leu Cys Arg Arg Ser Arg Ala Ala Ser Val Gly Arg Cys Ser Met Pro Glu Gly Pro Phe Pro Gly Arg Leu Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu

Val Cys Leu Thr Gly Gly Ser Glu Thr Ser Glu Phe Lys Phe Leu Lys
755 760 765

Pro Ile Ile Pro Asn Phe Ser Pro 770 775

<210> 442

<211> 776

<212> PRT

<213> Homo sapiens

<400> 442

Met Glu Ile Gly Trp Met His Asn Arg Arg Gln Arg Gln Val Leu Val

1 5 10 15

Phe Phe Val Leu Leu Ser Leu Ser Gly Ala Gly Ala Glu Leu Gly Ser 20 25 30

Tyr Ser Val Val Glu Glu Thr Glu Arg Gly Ser Phe Val Ala Asn Leu 35 40 45

Gly Lys Asp Leu Gly Leu Gly Leu Thr Glu Met Ser Thr Arg Lys Ala 50 55 60

Arg Ile Ile Ser Gln Gly Asn Lys Gln His Leu Gln Leu Lys Ala Gln 65 70 75 80

Thr Gly Asp Leu Leu Ile Asn Glu Lys Leu Asp Arg Glu Glu Leu Cys
85 90 95

Gly Pro Thr Glu Pro Cys Ile Leu His Phe Gln Val Leu Met Glu Asn 100 105 110

Pro Leu Glu Ile Phe Gln Ala Glu Leu Arg Val Ile Asp Ile Asn Asp 115 120 125

His Ser Pro Met Phe Thr Glu Lys Glu Met Ile Leu Lys Ile Pro Glu 130 135 140

Asn Ser Pro Leu Gly Thr Glu Phe Pro Leu Asn His Ala Leu Asp Leu 145 150 155 160

Asp Val Gly Ser Asn Asn Val Gln Asn Tyr Lys Ile Ser Pro Ser Ser 165 170 175

His Phe Arg Val Leu Ile His Glu Phe Arg Asp Gly Arg Lys Tyr Pro

| 180 | 185 | 190 |
|-----|-----|-----|
| | | |

- Glu Leu Val Leu Asp Lys Glu Leu Asp Arg Glu Glu Glu Pro Gln Leu 195 200 205

 Arg Leu Thr Leu Thr Ala Leu Asp Gly Gly Ser Pro Pro Arg Ser Gly
- 210 215 220
- Thr Ala Gln Val Arg Ile Glu Val Val Asp Ile Asn Asp Asn Ala Pro 225 230 235 240
- Glu Phe Glu Gln Pro Ile Tyr Lys Val Gln Ile Pro Glu Asn Ser Pro 245 250 255
- Leu Gly Ser Leu Val Ala Thr Val Ser Ala Arg Asp Leu Asp Gly Gly 260 265 270
- Ala Asn Gly Lys Ile Ser Tyr Thr Leu Phe Gln Pro Ser Glu Asp Ile 275 280 285
- Ser Lys Thr Leu Glu Val Asn Pro Met Thr Gly Glu Val Arg Leu Arg 290 295 300
- Lys Gln Val Asp Phe Glu Met Val Thr Ser Tyr Glu Val Arg Ile Lys 305 310 315 320
- Ala Thr Asp Gly Gly Leu Ser Gly Lys Cys Thr Leu Leu Gln 325 330 335
- Val Val Asp Val Asn Asp Asn Pro Pro Gln Val Thr Met Ser Ala Leu 340 345 350
- Thr Ser Pro Ile Pro Glu Asn Ser Pro Glu Ile Val Val Ala Val Phe 355 360 365
- Ser Val Ser Asp Pro Asp Ser Gly Asn Asn Gly Lys Thr Ile Ser Ser 370 380
- Ile Gln Glu Asp Leu Pro Phe Leu Leu Lys Pro Ser Val Lys Asn Phe 385 390 395 400
- Tyr Thr Leu Val Thr Glu Arg Ala Leu Asp Arg Glu Ala Arg Ala Glu 405 410 415
- Tyr Asn Ile Thr Leu Thr Val Thr Asp Met Gly Thr Pro Arg Leu Lys
 420 425 430
- Thr Glu His Asn Ile Thr Val Gln Ile Ser Asp Val Asn Asp Asn Ala

| 435 | 440 | 445 |
|-----|-----|-----|
| | | |

| Pro | Thr 450 | Phe | Thr | Gln | Thr | Ser 455 | Tyr | Thr | Leu | Phe | Val 460 | Arg | Glu | Asn | Asn |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Ser 465 | Pro | Ala | Leu | His | Ile 470 | Gly | Ser | Val | Ser | Ala 475 | Thr | Asp | Arg | Asp | Ser 480 |
| Gly | Thr | Asn | Ala | Gln 485 | Val | Thr | Tyr | Ser | Leu 490 | Leu | Pro | Pro | Gln | Asp 495 | Pro |
| His | Leu | Pro | Leu 500 | Ala | Ser | Leu | Val | Ser 505 | Ile | Asn | Ala | Asp | Asn 510 | Gly | His |
| Leu | Phe | Ala 515 | Leu | Arg | Ser | Leu | Asp 520 | Tyr | Glu | Ala | Leu | Arg 525 | Glu | Phe | Glu |
| Phe | Arg 530 | Val | Ser | Ala | Thr | Asp 535 | Arg | Gly | Ser | Pro | Ala 540 | Leu | Ser | Ser | Glu |
| Ala 545 | Leu | Val | Arg | Val | Leu 550 | Val | Leu | Asp | Ala | Asn 555 | Asp | Asn | Ser | Pro | Phe 560 |
| Val | Leu | Tyr | Pro | Leu 565 | Gln | Asn | Gly | Ser | Ala 570 | Pro | Cys | Thr | Glu | Leu 575 | Val |
| Pro | Arg | Ala | Ala 580 | Glu | Pro | Gly | Tyr | Leu 585 | Val | Thr | Lys | Val | Val 590 | Ala | Val |
| Asp | Gly | Asp 595 | Ser | Gly | Gln | Asn | Ala 600 | Trp | Leu | Ser | Tyr | Gln 605 | Leu | Leu | Lys |
| Ala | Thr 610 | Glu | Pro | Gly | Leu | Phe 615 | Gly | Val | Trp | Ala | His 620 | Asn | Gly | Glu | Val |
| Arg 625 | Thr | Ala | Arg | Leu | Leu 630 | Ser | Glu | Arg | Asp | Ala 635 | Ala | Lys | Gln | Arg | Leu 640 |
| Val | Val | Leu | Val | Lys 645 | Asp | Asn | Gly | Glu | Pro 650 | Pro | Arg | Ser | Ala | Thr 655 | Ala |
| Thr | Leu | His | Val 660 | Leu | Leu | Val | Asp | Gly 665 | Phe | Ser | Gln | Pro | Phe 670 | Leu | Pro |
| Leu | Pro | Glu 675 | Ala | Ala | Pro | Gly | Gln 680 | Thr | Gln | Ala | Asn | Ser 685 | Leu | Thr | Val |
| Tvr | Leu | Val | Val | Ala | Len | Ala | Ser | Val | Ser | Ser | Len | Phe | Len | Phe | Ser |

690 695 700

Val Leu Leu Phe Val Ala Val Arg Leu Cys Arg Arg Ser Arg Ala Ala 705 710 715 720

Ser Val Gly Arg Cys Ser Met Pro Glu Gly Pro Phe Pro Gly Arg Leu 725 730 735

Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu
740 745 750

Val Cys Leu Thr Gly Gly Ser Glu Thr Ser Glu Phe Lys Phe Leu Lys
755 760 765

Pro Ile Ile Pro Asn Phe Ser Pro 770 775

<210> 443

<211> 787

<212> PRT

<213> Homo sapiens

<400> 443

Ser Phe Cys Glu Pro Thr Phe Gln Glu Lys Ala Met Glu Ile Gly Trp
1 5 10 15

Met His Asn Arg Arg Gln Arg Gln Val Leu Val Phe Phe Val Leu Leu 20 25 30

Ser Leu Ser Gly Ala Gly Ala Glu Leu Gly Ser Tyr Ser Val Val Glu 35 40 45

Glu Thr Glu Arg Gly Ser Phe Val Ala Asn Leu Gly Lys Asp Leu Gly
50 55 60

Leu Gly Leu Thr Glu Met Ser Thr Arg Lys Ala Arg Ile Ile Ser Gln 65 70 75 80

Gly Asn Lys Gln His Leu Gln Leu Lys Ala Gln Thr Gly Asp Leu Leu 85 90 95

Ile Asn Glu Lys Leu Asp Arg Glu Glu Leu Cys Gly Pro Thr Glu Pro
100 105 110

Cys Ile Leu His Phe Gln Val Leu Met Glu Asn Pro Leu Glu Ile Phe 115 120 125

| Gln | Ala 130 | Glu | Leu | Arg | Val | Ile 135 | Asp | Ile | Asn | Asp | His 140 | Ser | Pro | Met | Phe |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Thr 145 | Glu | Lys | Glu | Met | Ile 150 | Leu | Lys | Ile | Pro | Glu 155 | Asn | Ser | Pro | Leu | Gly 160 |
| Thr | Glu | Phe | Pro | Leu 165 | Asn | His | Ala | Leu | Asp 170 | Leu | Asp | Val | Gly | Ser 175 | Asn |
| Asn | Val | Gln | Asn 180 | Tyr | Lys | Ile | Ser | Pro 185 | Ser | Ser | His | Phe | Arg 190 | Val | Leu |
| Ile | His | Glu 195 | Phe | Arg | Asp | Gly | Arg 200 | Lys | Tyr | Pro | Glu | Leu 205 | Val | Leu | Asp |
| Lys | Glu 210 | Leu | Asp | Arg | Glu | Glu 215 | Glu | Pro | Gln | Leu | Arg 220 | Leu | Thr | Leu | Thr |
| Ala 225 | Leu | Asp | Gly | Gly | Ser 230 | Pro | Pro | Arg | Ser | Gly 235 | Thr | Ala | Gln | Val | Arg 240 |
| Ile | Glu | Val | Val | Asp 245 | Ile | Asn | Asp | Asn | Ala 250 | Pro | Glu | Phe | Glu | Gln 255 | Pro |
| Ile | Tyr | Lys | Val 260 | Gln | Ile | Pro | Glu | Asn 265 | Ser | Pro | Leu | Gly | Ser 270 | Leu | Val |
| Ala | Thr | Val 275 | Ser | Ala | Arg | Asp | Leu 280 | Asp | Gly | Gly | Ala | Asn 285 | Gly | Lys | Ile |
| Ser | Tyr 290 | Thr | Leu | Phe | Gln | Pro 295 | Ser | Glu | Asp | Ile | Ser 300 | Lys | Thr | Leu | Glu |
| Val 305 | Asn | Pro | Met | Thr | Gly 310 | Glu | Val | Arg | Leu | Arg 315 | Lys | Gln | Val | Asp | Phe 320 |
| Glu | Met | Val | Thr | Ser 325 | Tyr | Glu | Val | Arg | Ile 330 | Lys | Ala | Thr | Asp | Gly 335 | Gly |
| Gly | Leu | Ser | Gly 340 | Lys | Cys | Thr | Leu | Leu 345 | Leu | Gln | Val | Val | Asp 350 | Val | Asn |
| Asp | Asn | Pro 355 | Pro | Gln | Val | Thr | Met 360 | Ser | Ala | Leu | Thr | Ser 365 | Pro | Ile | Pro |
| Glu | Asn 370 | Ser | Pro | Glu | Ile | Val 375 | Val | Ala | Val | Phe | Ser 380 | Val | Ser | Asp | Pro |

| Asp 385 | Ser | Gly | Asn | Asn | Gly 390 | Lys | Thr | Ile | Ser | Ser 395 | Ile | Gln | Glu | Asp | Leu 400 |
|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|------------|
| Pro | Phe | Leu | Leu | Lys 405 | Pro | Ser | Val | Lys | Asn 410 | Phe | Tyr | Thr | Leu | Val 415 | Thr |
| Glu | Arg | Ala | Leu 420 | Asp | Arg | Glu | Ala | Arg 425 | Ala | Glu | Tyr | Asn | Ile 430 | Thr | Leu |
| Thr | Val | Thr 435 | Asp | Met | Gly | Thr | Pro 440 | Arg | Leu | Lys | Thr | Glu 445 | His | Asn | Ile |
| Thr | Val 450 | Gln | Ile | Ser | Asp | Val 455 | Asn | Asp | Asn | Ala | Pro 460 | Thr | Phe | Thr | Gln |
| Thr 465 | Ser | Tyr | Thr | Leu | Phe 470 | Val | Arg | Glu | Asn | Asn 475 | Ser | Pro | Ala | Leu | His 480 |
| Ile | Gly | Ser | Val | Ser 485 | Ala | Thr | Asp | Arg | Asp 490 | Ser | Gly | Ile | Asn | Ala 495 | Gln |
| Val | Thr | Tyr | Ser 500 | Leu | Leu | Pro | Pro | Gln 505 | Asp | Pro | His | Leu | Pro 510 | Leu | Ala |
| Ser | Leu | Val 515 | Ser | Ile | Asn | Ala | Asp 520 | Asn | Gly | His | Leu | Phe 525 | Ala | Leu | Arg |
| Ser | Leu 530 | Asp | Tyr | Glu | Ala | Leu 535 | Arg | Glu | Phe | Glu | Phe 540 | Arg | Val | Ser | Ala |
| Thr 545 | Asp | Arg | Gly | Ser | Pro 550 | Ala | Leu | Ser | Ser | Glu 555 | Ala | Leu | Val | Arg | Val 560 |
| Leu | Val | Leu | Asp | Ala 565 | Asn | Asp | Asn | Ser | Pro 570 | Phe | Val | Leu | Tyr | Pro 575 | Leu |
| Gln | Asn | Gly | Ser 580 | Ala | Pro | Cys | Thr | Glu 585 | Leu | Val | Pro | Arg | Ala 590 | Ala | Glu |
| Pro | Gly | Tyr 595 | Leu | Val | Thr | Lys | Val 600 | Val | Ala | Val | Asp | Gly 605 | Asp | Ser | Gly |
| Gln | Asn 610 | Ala | Trp | Leu | Ser | Tyr 615 | Gln | Leu | Leu | Lys | Ala 620 | Thr | Glu | Pro | Gly |
| Leu 625 | Phe | Gly | Val | Trp | Ala 630 | His | Asn | Gly | Glu | Val 635 | Arg | Thr | Ala | Arg | Leu 640 |

. .

Leu Ser Glu Arg Asp Ala Ala Lys His Arg Leu Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Cys Ser Ala Thr Ala Thr Leu His Val Leu Leu Val Asp Gly Phe Ser Gln Pro Phe Leu Pro Leu Pro Glu Ala Ala Pro Gly Gln Thr Gln Ala Asn Ser Leu Thr Val Tyr Leu Val Val Ala Leu Ala Ser Val Ser Ser Leu Phe Leu Phe Ser Val Leu Leu Phe Val Val Val Arg Leu Cys Arg Arg Ser Arg Ala Ala Ser Val Gly Arg Cys Ser Met Pro Glu Gly Pro Phe Pro Gly Arg Leu Val Asp Val Ser Gly Thr Gly Thr Leu Ser Gln Ser Tyr Gln Tyr Glu Val Cys Leu Thr Gly Gly Ser Glu Thr Ser Glu Phe Lys Phe Leu Lys Pro Ile Ile Pro Asn Phe Ser Pro <210> 444 <211> 87 <212> PRT <213> Homo sapiens <400> 444 Val Ser Ala Thr Asp Arg Asp Ser Gly Thr Asn Ala Gln Val Thr Tyr Ser Leu Leu Pro Pro Gln Asp Pro His Leu Pro Leu Ser Ser Leu Val Ser Ile Asn Ala Asp Asn Gly His Leu Phe Ala Leu Arg Ser Leu Asp Tyr Glu Ala Leu Gln Ala Phe Glu Phe Arg Val Gly Ala Thr Asp Arg

Gly Ser Pro Ala Leu Ser Ser Glu Ala Leu Val Arg Val Leu Val Leu 75 65 70 Asp Ala Asn Asp Asn Ser Pro 85 <210> 445 <211> 82 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Cadherin repeats domain sequence <400> 445 Val Ser Ala Thr Asp Ala Asp Ser Gly Glu Asn Gly Lys Val Thr Tyr 1 10 15 Ser Ile Leu Ser Gly Asn Asp Gly Gly Leu Phe Ser Ile Asp Pro Glu 20 25 Thr Gly Ile Ile Thr Thr Lys Pro Leu Asp Arg Glu Glu Gln Ser Glu Tyr Thr Leu Thr Val Glu Ala Thr Asp Gly Gly Pro Pro Leu 55 Ser Ser Thr Ala Thr Val Thr Val Leu Asp Val Asn Asp Asn 75 65 70 Ala Pro <210> 446 <211> 82 <212> PRT <213> Homo sapiens <400> 446 Val Ser Ala Arg Asp Leu Asp Ile Gly Thr Asn Gly Glu Ile Ser Tyr 5 10 15 1 Ala Phe Ser Gln Ala Ser Glu Asp Ile Arg Lys Thr Phe Arg Leu Ser

20

25

Ala Lys Ser Gly Glu Leu Leu Leu Arg Gln Lys Leu Asp Phe Glu Ser 35 40 45

Ile Gln Thr Tyr Thr Val Asn Ile Gln Ala Thr Asp Gly Gly Leu 50 55 60

Ser Gly Lys Ser Thr Val Ile Val Gln Val Val Asp Val Asn Asp Asn 65 70 75 80

Pro Pro

<210> 447

<211> 85

<212> PRT

<213> Homo sapiens

<400> 447

Asn Ala Glu Asp Leu Asp Val Gly Arg Asn Ser Leu Gln Asn Tyr Thr

1 5 10 15

Ile Thr Pro Asn Ser His Phe His Val Pro Thr Arg Ser Arg Arg Asp
20 25 30

Gly Arg Lys Tyr Pro Glu Leu Val Leu Asn Arg Ala Leu Asp Arg Glu 35 40 45

Glu Gln Pro Glu Ile Arg Leu Thr Leu Thr Ala Leu Asp Gly Gly Ser
50 55 60

Pro Pro Arg Ser Gly Thr Ala Leu Val Arg Ile Glu Val Val Asp Ile 65 70 75 80

Asn Asp Asn Val Pro

85

<210> 448

<211> 81

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cadherin repeats domain sequence

<400> 448

Ser Ala Thr Asp Ala Asp Ser Gly Glu Asn Gly Lys Val Thr Tyr Ser 1 5 10 15

Ile Leu Ser Gly Asn Asp Gly Gly Leu Phe Ser Ile Asp Pro Glu Thr
20 25 30

Gly Ile Ile Thr Thr Lys Pro Leu Asp Arg Glu Glu Gln Ser Glu 35 40 45

Tyr Thr Leu Thr Val Glu Ala Thr Asp Gly Gly Pro Pro Leu Ser 50 55 60

Ser Thr Ala Thr Val Thr Val Leu Asp Val Asn Asp Asn Ala 65 70 75 80

Pro

<210> 449

<211> 81

<212> PRT

<213> Homo sapiens

<400> 449

Ser Val Ser Asp Leu Asp Ser Gly Asp Asn Gly Arg Val Met Cys Ser 1 5 10 15

Ile Glu Asn Asn Leu Pro Phe Phe Leu Lys Pro Ser Val Glu Asn Phe 20 25 30

Tyr Thr Leu Val Ser Glu Gly Ala Leu Asp Arg Glu Thr Arg Ser Glu
35 40 45

Tyr Asn Ile Thr Ile Thr Ile Thr Asp Leu Gly Thr Pro Arg Leu Lys
50 55 60

Thr Lys Tyr Asn Ile Thr Val Leu Val Ser Asp Val Asn Asp Asn Ala 65 70 75 80

Pro

<210> 450

<211> 76

<212> PRT

<213> Homo sapiens <400> 450 Val Val Ala Val Asp Gly Asp Ser Gly Gln Asn Ala Trp Leu Ser Tyr 5 10 1 15 Gln Leu Leu Lys Ala Thr Glu Pro Gly Leu Phe Gly Val Trp Ala His 20 25 30 Asn Gly Glu Val Arg Thr Ala Arg Leu Leu Arg Glu Arg Asp Ala Ala 35 45 Lys Gln Arg Leu Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg 50 55 60 Ser Ala Thr Ala Thr Leu His Val Leu Leu Val Asp 70 75 65 <210> 451 <211> 76 <212> PRT <213> Artificial Sequence <220> <223> Description of Artificial Sequence: Cadherin repeats domain sequence <400> 451 Val Ser Ala Thr Asp Ala Asp Ser Gly Glu Asn Gly Lys Val Thr Tyr 1 10 15 Ser Ile Leu Ser Gly Asn Asp Gly Gly Leu Phe Ser Ile Asp Pro Glu 20 25

Thr Gly Ile Ile Thr Thr Lys Pro Leu Asp Arg Glu Glu Gln Ser 35 40 45

Glu Tyr Thr Leu Thr Val Glu Ala Thr Asp Gly Gly Gro Pro Leu 55

Ser Ser Thr Ala Thr Val Thr Val Leu Asp 65 70 75

<210> 452 <211> 91 <212> PRT

<213> Homo sapiens

<400> 452

Tyr Glu Val Gln Ile Pro Glu Asp Ser Pro Val Gly Ser Gln Val Ala 1 5 10 15

Ile Val Ser Ala Arg Asp Leu Asp Ile Gly Thr Asn Gly Glu Ile Ser 20 25 30

Tyr Ala Phe Ser Gln Ala Ser Glu Asp Ile Arg Lys Thr Phe Arg Leu 35 40 45

Ser Ala Lys Ser Gly Glu Leu Leu Leu Arg Gln Lys Leu Asp Phe Glu 50 55 60

Ser Ile Gln Thr Tyr Thr Val Asn Ile Gln Ala Thr Asp Gly Gly Gly 65 70 75 80

Leu Ser Gly Lys Ser Thr Val Ile Val Gln Val 85 90

<210> 453

<211> 91

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cadherin repeats domain sequence

<400> 453

Tyr Ser Ala Ser Val Pro Glu Asn Ala Pro Val Gly Thr Glu Val Leu
1 5 10 15

Thr Val Thr Ala Thr Asp Ala Asp Leu Gly Pro Asn Gly Arg Ile Phe 20 25 30

Tyr Ser Ile Leu Gly Gly Gly Pro Gly Gly Trp Phe Arg Ile Asp Pro
35 40 45

Asp Thr Gly Asp Leu Ser Thr Thr Lys Pro Leu Asp Arg Glu Ser Ile 50 55 60

Gly Glu Tyr Glu Leu Thr Val Leu Ala Thr Asp Ser Gly Gly Pro Pro 65 70 75 80

Leu Ser Gly Thr Thr Thr Val Thr Ile Thr Val

85 90

<210> 454

<211> 97

<212> PRT

<213> Homo sapiens

<400> 454

Tyr Thr Leu Phe Val Arg Glu Asn Asn Ser Pro Ala Leu His Ile Gly
1 5 10 15

Ser Val Ser Ala Thr Asp Arg Asp Ser Gly Thr Asn Ala Gln Val Thr 20 25 30

Tyr Ser Leu Leu Pro Pro Gln Asp Pro His Leu Pro Leu Ser Ser Leu 35 40 45

Val Ser Ile Asn Ala Asp Asn Gly His Leu Phe Ala Leu Arg Ser Leu 50 55 60

Asp Tyr Glu Ala Leu Gln Ala Phe Glu Phe Arg Val Gly Ala Thr Asp 65 70 75 80

Arg Gly Ser Pro Ala Leu Ser Ser Glu Ala Leu Val Arg Val Leu Val 85 90 95

Leu

<210> 455

<211> 92

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cadherin repeats domain sequence

<400> 455

Tyr Ser Ala Ser Val Pro Glu Asn Ala Pro Val Gly Thr Glu Val Leu
1 5 10 15

Thr Val Thr Ala Thr Asp Ala Asp Leu Gly Pro Asn Gly Arg Ile Phe 20 . 25 . 30

Tyr Ser Ile Leu Gly Gly Gly Pro Gly Gly Trp Phe Arg Ile Asp Pro

35 40 45

Asp Thr Gly Asp Leu Ser Thr Thr Lys Pro Leu Asp Arg Glu Ser Ile 50 55 60

Gly Glu Tyr Glu Leu Thr Val Leu Ala Thr Asp Ser Gly Gly Pro Pro 65 70 75 80

Leu Ser Gly Thr Thr Thr Val Thr Ile Thr Val Leu 85 90

<210> 456

<211> 85

<212> PRT

<213> Homo sapiens

<400> 456

Val Pro Arg Ala Ala Glu Pro Gly Tyr Leu Val Thr Lys Val Val Ala 1 5 10 15

Val Asp Gly Asp Ser Gly Gln Asn Ala Trp Leu Ser Tyr Gln Leu Leu 20 25 30

Lys Ala Thr Glu Pro Gly Leu Phe Gly Val Trp Ala His Asn Gly Glu $35 \hspace{1cm} 40 \hspace{1cm} 45$

Val Arg Thr Ala Arg Leu Leu Arg Glu Arg Asp Ala Ala Lys Gln Arg 50 55 \cdot 60

Leu Val Val Leu Val Lys Asp Asn Gly Glu Pro Pro Arg Ser Ala Thr 65 70 75 80

Ala Thr Leu His Val

85

<210> 457

<211> 85

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cadherin repeats domain sequence

<400> 457

Val Pro Glu Asn Ala Pro Val Gly Thr Glu Val Leu Thr Val Thr Ala

Thr Asp Ala Asp Leu Gly Pro Asn Gly Arg Ile Phe Tyr Ser Ile Leu 20 25 30

Gly Gly Pro Gly Gly Trp Phe Arg Ile Asp Pro Asp Thr Gly Asp
35 40 45

Leu Ser Thr Thr Lys Pro Leu Asp Arg Glu Ser Ile Gly Glu Tyr Glu
50 55 60

Leu Thr Val Leu Ala Thr Asp Ser Gly Gly Pro Pro Leu Ser Gly Thr
65 70 75 80

Thr Thr Val Thr Ile

<210> 458

1

<211> 91

<212> PRT

<213> Homo sapiens

<400> 458

Ile Leu Glu Ser Thr Leu Pro Gly Thr Val Ile Pro Leu Gly Asn Ala 1 5 10 15

Glu Asp Leu Asp Val Gly Arg Asn Ser Leu Gln Asn Tyr Thr Ile Thr
20 25 30

Pro Asn Ser His Phe His Val Pro Thr Arg Ser Arg Asp Gly Arg 35 40 45

Lys Tyr Pro Glu Leu Val Leu Asn Arg Ala Leu Asp Arg Glu Glu Gln 50 55 60

Pro Glu Ile Arg Leu Thr Leu Thr Ala Leu Asp Gly Gly Ser Pro Pro 65 70 75 80

Arg Ser Gly Thr Ala Leu Val Arg Ile Glu Val 85 90

<210> 459

<211> 87

<212> PRT

<213> Artificial Sequence

<220>

<223> Description of Artificial Sequence: Cadherin repeats domain sequence

<400> 459

Val Pro Glu Asn Ala Pro Val Gly Thr Glu Val Leu Thr Val Thr Ala
1 5 10 15

Thr Asp Ala Asp Leu Gly Pro Asn Gly Arg Ile Phe Tyr Ser Ile Leu 20 25 30

Gly Gly Gly Pro Gly Gly Trp Phe Arg Ile Asp Pro Asp Thr Gly Asp 35 40 45

Leu Ser Thr Thr Lys Pro Leu Asp Arg Glu Ser Ile Gly Glu Tyr Glu
50 55 60

Leu Thr Val Leu Ala Thr Asp Ser Gly Gly Pro Pro Leu Ser Gly Thr 65 70 75 80

Thr Thr Val Thr Ile Thr Val 85

<210> 460

<211> 86

<212> PRT

<213> Homo sapiens

<400> 460

Asp Leu Asp Ser Gly Asp Asn Gly Arg Val Met Cys Ser Ile Glu Asn 20 25 30

Asn Leu Pro Phe Phe Leu Lys Pro Ser Val Glu Asn Phe Tyr Thr Leu
35 40 45

Val Ser Glu Gly Ala Leu Asp Arg Glu Thr Arg Ser Glu Tyr Asn Ile 50 55 60

Thr Ile Thr Ile Thr Asp Leu Gly Thr Pro Arg Leu Lys Thr Lys Tyr
65 70 75 80

Asn Ile Thr Val Leu Val

85

| | 461 | | | |
|--|--|-----------|------------|----|
| <211> | 22 | | | |
| <212> | DNA | | | |
| <213> | Artificial Sequence | | | |
| | - | | | |
| <220> | | | | |
| | Description of Artificial | Sequence: | PCR Primer | |
| | sequence | | | |
| | 0.4.0 | | | |
| <400> | 461 | | | |
| | itett tggetgtgaa gt | | | 22 |
| ~~5~5 | | | | |
| | | | | |
| <210> | 462 | | | |
| <211> | | | | |
| <212> | | | | |
| | Artificial Sequence | | | |
| \213/ | Artificial bequeince | | | |
| <220> | | | | |
| | Description of Artificial | Cognongo | DCD Drimon | |
| \ 2237 | Description of Artificial | sequence: | PCK PIIMEI | |
| | sequence | | | |
| < 4.0.0> | 462 | | | |
| <400> | | | | 22 |
| ctacco | ccatg gcctccatcg agt | | | 23 |
| | | | | |
| | | | | |
| <010> | 463 | | | |
| <210> | | | | |
| <211> | 19 | | | |
| <211> <212> | 19 DNA | | | |
| <211> <212> | 19 | | | |
| <211> <212> <213> | 19 DNA | | | |
| <211> <212> <213> <220> | 19 DNA Artificial Sequence | | | |
| <211> <212> <213> <220> | 19 DNA Artificial Sequence Description of Artificial | Sequence: | PCR Primer | |
| <211> <212> <213> <220> | 19 DNA Artificial Sequence | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> | 19 DNA Artificial Sequence Description of Artificial sequence | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> <400> | 19 DNA Artificial Sequence Description of Artificial sequence 463 | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> <400> | 19 DNA Artificial Sequence Description of Artificial sequence | Sequence: | PCR Primer | 19 |
| <211> <212> <213> <220> <223> <400> | 19 DNA Artificial Sequence Description of Artificial sequence 463 | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> <400> ggatgt | Description of Artificial sequence 463 ccaa gccatcctt | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> <400> ggatgt | Description of Artificial sequence 463 ccaa gccatcctt | Sequence: | PCR Primer | |
| <211> <212> <213> <220> <223> <400> ggatgt <210> <211> | Description of Artificial sequence 463 ccaa gccatcctt | Sequence: | PCR Primer | |
| <211><212><213> 220 223 400 <td>Description of Artificial sequence 463 ccaa gccatcctt 464 22 DNA</td> <td>Sequence:</td> <td>PCR Primer</td> <td></td> | Description of Artificial sequence 463 ccaa gccatcctt 464 22 DNA | Sequence: | PCR Primer | |
| <211><212><213> 220 223 400 <td>Description of Artificial sequence 463 ccaa gccatcctt</td> <td>Sequence:</td> <td>PCR Primer</td> <td></td> | Description of Artificial sequence 463 ccaa gccatcctt | Sequence: | PCR Primer | |
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